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1 *Streptococcus pneumoniae* Antigens and Vaccines



Field of the Invention

The present invention relates to novel *Streptococcus pneumoniae* antigens for the detection of *Streptococcus* and for the prevention or attenuation of disease caused by *Streptococcus*. The invention further relates to isolated nucleic acid molecules encoding antigenic polypeptides of *S. pneumoniae*. Antigenic polypeptides are also provided, as are vectors, host cells and recombinant methods for producing the same. The invention additionally relates to diagnostic methods for detecting *Streptococcus* gene expression.

Background of the Invention

Streptococcus pneumoniae has been one of the most extensively studied microorganisms since its first isolation in 1881. It was the object of many investigations that led to important scientific discoveries. In 1928, Griffith observed that when heat-killed encapsulated pneumococci and live strains constitutively lacking any capsule were concomitantly injected into mice, the nonencapsulated could be converted into encapsulated pneumococci with the same capsular type as the heat-killed strain. Years later, the nature of this "transforming principle," or carrier of genetic information, was shown to be DNA. (Avery, O.T., *et al.*, *J. Exp. Med.*, 79:137-157 (1944)).

In spite of the vast number of publications on *S. pneumoniae* many questions about its virulence are still unanswered, and this pathogen remains a major causative agent of serious human disease, especially community-acquired pneumonia. (Johnston, R.B., *et al.*, *Rev. Infect. Dis.* 13(Suppl. 6):S509-517 (1991)). In addition, in developing countries, the pneumococcus is responsible for the death of a large number of children under the age of 5 years from pneumococcal pneumonia. The incidence of pneumococcal disease is highest in infants under 2 years of age and in people over 60 years of age. Pneumococci are the second most frequent cause (after *Haemophilus influenzae* type b) of bacterial meningitis and otitis media in children. With the recent introduction of conjugate vaccines for *H. influenzae* type b, pneumococcal meningitis is likely to become increasingly prominent. *S. pneumoniae* is the most important etiologic agent of community-acquired pneumonia in adults and is the second most common cause of bacterial meningitis behind *Neisseria meningitidis*.

The antibiotic generally prescribed to treat *S. pneumoniae* is benzylpenicillin, although resistance to this and to other antibiotics is found occasionally. Pneumococcal resistance to penicillin results from mutations in its

penicillin-binding proteins. In uncomplicated pneumococcal pneumonia caused by a sensitive strain, treatment with penicillin is usually successful unless started too late. Erythromycin or clindamycin can be used to treat pneumonia in patients hypersensitive to penicillin, but resistant strains to these drugs exist. Broad spectrum antibiotics (e.g., the tetracyclines) may also be effective, although tetracycline-resistant strains are not rare. In spite of the availability of antibiotics, the mortality of pneumococcal bacteremia in the last four decades has remained stable between 25 and 29%. (Gillespie, S.H., *et al.*, *J. Med. Microbiol.* 28:237-248 (1989).

S. pneumoniae is carried in the upper respiratory tract by many healthy individuals. It has been suggested that attachment of pneumococci is mediated by a disaccharide receptor on fibronectin, present on human pharyngeal epithelial cells. (Anderson, B.J., *et al.*, *J. Immunol.* 142:2464-2468 (1989). The mechanisms by which pneumococci translocate from the nasopharynx to the lung, thereby causing pneumonia, or migrate to the blood, giving rise to bacteremia or septicemia, are poorly understood. (Johnston, R.B., *et al.*, *Rev. Infect. Dis.* 13(Suppl. 6):S509-517 (1991).

Various proteins have been suggested to be involved in the pathogenicity of *S. pneumoniae*, however, only a few of them have actually been confirmed as virulence factors. Pneumococci produce an IgA1 protease that might interfere with host defense at mucosal surfaces. (Kornfield, S.J., *et al.*, *Rev. Inf. Dis.* 3:521-534 (1981). *S. pneumoniae* also produces neuraminidase, an enzyme that may facilitate attachment to epithelial cells by cleaving sialic acid from the host glycolipids and gangliosides. Partially purified neuraminidase was observed to induce meningitis-like symptoms in mice; however, the reliability of this finding has been questioned because the neuraminidase preparations used were probably contaminated with cell wall products. Other pneumococcal proteins besides neuraminidase are involved in the adhesion of pneumococci to epithelial and endothelial cells. These pneumococcal proteins have as yet not been identified. Recently, Cundell *et al.*, reported that peptide permeases can modulate pneumococcal adherence to epithelial and endothelial cells. It was, however, unclear whether these permeases function directly as adhesions or whether they enhance adherence by modulating the expression of pneumococcal adhesions. (DeVelasco, E.A., *et al.*, *Micro. Rev.* 59:591-603 (1995). A better understanding of the virulence factors determining its pathogenicity will need to be developed to cope with the devastating effects of pneumococcal disease in humans.

Ironically, despite the prominent role of *S. pneumoniae* in the discovery of DNA, little is known about the molecular genetics of the organism. The *S. pneumoniae* genome consists of one circular, covalently closed, double-stranded DNA and a collection of so-called variable accessory elements, such as prophages, plasmids, transposons and the like. Most physical characteristics and almost all of the genes of *S. pneumoniae* are unknown. Among the few that have been identified, most have not been physically mapped or characterized in detail. Only a few genes of this organism have been sequenced. (See, for instance current versions of GENBANK and other nucleic acid databases, and references that relate to the genome of *S. pneumoniae* such as those set out elsewhere herein.) Identification of *in vivo*-expressed, and broadly protective, antigens of *S. pneumoniae* has remained elusive.

Summary of the Invention

The present invention provides isolated nucleic acid molecules comprising polynucleotides encoding the *S. pneumoniae* polypeptides described in Table 1 and having the amino acid sequences shown as ~~SEQ ID NO: 2, SEQ ID NO: 4, SEQ ID NO: 6, and so on through SEQ ID NO: 226~~. Thus, one aspect of the invention provides isolated nucleic acid molecules comprising polynucleotides having a nucleotide sequence selected from the group consisting of: (a) a nucleotide sequence encoding ~~any of~~ the amino acid sequences of the polypeptides shown in Table 1; and (b) a nucleotide sequence complementary to any of the nucleotide sequences in (a).

Further embodiments of the invention include isolated nucleic acid molecules that comprise a polynucleotide having a nucleotide sequence at least 90% identical, and more preferably at least 95%, 96%, 97%, 98% or 99% identical, to any of the nucleotide sequences in (a) or (b) above, or a polynucleotide which hybridizes under stringent hybridization conditions to a polynucleotide in (a) or (b) above. This polynucleotide which hybridizes does not hybridize under stringent hybridization conditions to a polynucleotide having a nucleotide sequence consisting of only A residues or of only T residues. Additional nucleic acid embodiments of the invention relate to isolated nucleic acid molecules comprising polynucleotides which encode the amino acid sequences of epitope-bearing portions of an *S. pneumoniae* polypeptide having an amino acid sequence in (a) above.

The present invention also relates to recombinant vectors, which include the isolated nucleic acid molecules of the present invention, and to host cells containing the recombinant vectors, as well as to methods of making such

vectors and host cells and for using these vectors for the production of *S. pneumoniae* polypeptides or peptides by recombinant techniques.

The invention further provides isolated *S. pneumoniae* polypeptides having an amino acid sequence selected from the group consisting of an amino acid sequence of ~~any of~~ the polypeptides described in Table 1.

The polypeptides of the present invention also include polypeptides having an amino acid sequence with at least 70% similarity, and more preferably at least 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% similarity to those described in Table 1, as well as polypeptides having an amino acid sequence at least 70% identical, more preferably at least 75% identical, and still more preferably 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% identical to those above; as well as isolated nucleic acid molecules encoding such polypeptides.

The present invention further provides a vaccine, preferably a multi-component vaccine comprising one or more of the *S. pneumoniae* polynucleotides or polypeptides described in Table 1, or fragments thereof, together with a pharmaceutically acceptable diluent, carrier, or excipient, wherein the *S. pneumoniae* polypeptide(s) are present in an amount effective to elicit an immune response to members of the *Streptococcus* genus in an animal. The *S. pneumoniae* polypeptides of the present invention may further be combined with one or more immunogens of one or more other streptococcal or non-streptococcal organisms to produce a multi-component vaccine intended to elicit an immunological response against members of the *Streptococcus* genus and, optionally, one or more non-streptococcal organisms.

The vaccines of the present invention can be administered in a DNA form, e.g., "naked" DNA, wherein the DNA encodes one or more streptococcal polypeptides and, optionally, one or more polypeptides of a non-streptococcal organism. The DNA encoding one or more polypeptides may be constructed such that these polypeptides are expressed fusion proteins.

The vaccines of the present invention may also be administered as a component of a genetically engineered organism. Thus, a genetically engineered organism which expresses one or more *S. pneumoniae* polypeptides may be administered to an animal. For example, such a genetically engineered organism may contain one or more *S. pneumoniae* polypeptides of the present invention intracellularly, on its cell surface, or in its periplasmic space. Further, such a genetically engineered organism may secrete one or more *S. pneumoniae* polypeptides.

The vaccines of the present invention may be co-administered to an animal with an immune system modulator (e.g., CD86 and GM-CSF).

The invention also provides a method of inducing an immunological response in an animal to one or more members of the *Streptococcus* genus, ^{preferably} one or more isolates of the *S. pneumoniae* genus, comprising administering to the animal a vaccine as described above.

The invention further provides a method of inducing a protective immune response in an animal, sufficient to prevent or attenuate an infection by members of the *Streptococcus* genus, preferably at least *S. pneumoniae*, comprising administering to the animal a composition comprising ~~one or more~~ of the polynucleotides or polypeptides described in Table 1, or fragments thereof. Further, these polypeptides, or fragments thereof, may be conjugated to another immunogen and/or administered in admixture with an adjuvant.

The invention further relates to antibodies elicited in an animal by the administration of one or more *S. pneumoniae* polypeptides of the present invention and to methods for producing such antibodies.

The invention also provides diagnostic methods for detecting the expression of genes of members of the *Streptococcus* genus in an animal. One such method involves assaying for the expression of a gene encoding *S. pneumoniae* peptides in a sample from an animal. This expression may be assayed either directly (e.g., by assaying polypeptide levels using antibodies elicited in response to amino acid sequences described in Table 1) or indirectly (e.g., by assaying for antibodies having specificity for ^{the} amino acid sequences described in Table 1). An example of such a method involves the use of the polymerase chain reaction (PCR) to amplify and detect *Streptococcus* nucleic acid sequences.

The present invention also relates to nucleic acid probes having all or part of a nucleotide sequence described in Table 1 (shown as ^{SEQ ID NO:1} ~~SEQ ID NO:1, SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225~~ ^{which is} ~~which are~~ capable of hybridizing under stringent conditions to *Streptococcus* nucleic acids. The invention further relates to a method of detecting one or more *Streptococcus* nucleic acids in a biological sample obtained from an animal, said one or more nucleic acids encoding *Streptococcus* polypeptides, comprising: (a) contacting the sample with one or more of the above-described nucleic acid probes, under conditions such that hybridization occurs, and (b) detecting hybridization of said one or more probes to the *Streptococcus* nucleic acid present in the biological sample.

The invention also includes immunoassays, including an immunoassay for detecting *Streptococcus*, preferably at least isolates of the *S. pneumoniae* genus, comprising incubation of a sample (which is suspected of being infected with *Streptococcus*) with a probe antibody directed against an antigen/epitope of *S. pneumoniae*, to be detected under conditions allowing the formation of an antigen-antibody complex; and detecting the antigen-antibody complex which contains the probe antibody. An immunoassay for the detection of antibodies which are directed against a *Streptococcus* antigen comprising the incubation of a sample (containing antibodies from a mammal suspected of being infected with *Streptococcus*) with a probe polypeptide including an epitope of *S. pneumoniae*, under conditions that allow the formation of antigen-antibody complexes which contain the probe epitope containing antigen.

Some aspects of the invention pertaining to kits are those for: investigating samples for the presence of polynucleotides derived from *Streptococcus* which comprise a polynucleotide probe including a nucleotide sequence selected from Table 1 or a fragment thereof of approximately 15 or more nucleotides, in an appropriate container; analyzing the samples for the presence of antibodies directed against a *Streptococcus* antigen made up of a polypeptide which contains a *S. pneumoniae* epitope present in the polypeptide, in a suitable container; and analyzing samples for the presence of *Streptococcus* antigens made up of an anti-*S. pneumoniae* antibody, in a suitable container.

Detailed Description

The present invention relates to recombinant antigenic *S. pneumoniae* polypeptides and fragments thereof. The invention also relates to methods for using these polypeptides to produce immunological responses and to confer immunological protection to disease caused by members of the genus *Streptococcus*, at least isolates of the *S. pneumoniae* genus. The invention further relates to nucleic acid sequences which encode antigenic *S. pneumoniae* polypeptides and to methods for detecting *S. pneumoniae* nucleic acids and polypeptides in biological samples. The invention also relates to *S. pneumoniae*-specific antibodies and methods for detecting such antibodies produced in a host animal.

Definitions

The following definitions are provided to clarify the subject matter which the inventors consider to be the present invention.

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As used herein, the phrase "pathogenic agent" means an agent which causes a disease state or affliction in an animal. Included within this definition, for examples, are bacteria, protozoans, fungi, viruses and metazoan parasites which either produce a disease state or render an animal infected with such an organism susceptible to a disease state (*e.g.*, a secondary infection). Further included are species and strains of the genus *Streptococcus* which produce disease states in animals.

As used herein, the term "organism" means any living biological system, including viruses, regardless of whether it is a pathogenic agent.

As used herein, the term "*Streptococcus*" means any species or strain of bacteria which is members of the genus *Streptococcus*. Such species and strains are known to those of skill in the art, and include those that are pathogenic and those that are not.

As used herein, the phrase "one or more *S. pneumoniae* polypeptides of the present invention" means polypeptides comprising the amino acid sequence ~~of one or more of~~ the *S. pneumoniae* polypeptides described in Table 1 and disclosed as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226. These polypeptides may be expressed as fusion proteins wherein the *S. pneumoniae* polypeptides of the present invention are linked to additional amino acid sequences which may be of streptococcal or non-streptococcal origin. This phrase further includes polypeptide comprising fragments of the *S. pneumoniae* polypeptides of the present invention.

Additional definitions are provided throughout the specification.

Explanation of Table 1

Table 1, below, provides information describing ^{one} ~~the~~ open reading frames (ORFs) which encode potentially antigenic polypeptides of *S. pneumoniae* of the present invention. The table lists the ORF identifier which consists of the letters SP, which denote *S. pneumoniae*, followed immediately by a three digit numeric code, which arbitrarily number the potentially antigenic polypeptides of *S. pneumoniae* of the present invention and the nucleotide or amino acid sequence of ^{the} ~~each~~ ORF and encoded polypeptide. The table further correlates the ORF identifier with a sequence identification number (SEQ ID NO:). The actual nucleotide or amino acid sequence of ^{the} ~~each~~ ORF identifier is also shown in the Sequence Listing under the corresponding SEQ ID NO.

Thus, for example, the designation "SP⁰³⁶₁₂₆" refers to both the nucleotide and amino acid sequences of *S. pneumoniae* polypeptide number ⁰³⁶ ~~126~~ of the present invention. Further, "SP⁰³⁶₁₂₆" correlates with the nucleotide

sequence shown as SEQ ID NO: ~~223~~²²⁴ and with the amino acid sequence shown as SEQ ID NO: ~~224~~²²⁵ as is described in Table 1.

The open reading frame within each "ORF" begins with the second nucleotide shown. Thus, the first codon for each nucleotide sequence shown is bases 2-4, the second 5-7, the third 8-10, and so on.

Explanation of Table 2

Table 2 lists the antigenic epitopes present in ~~each of~~ the *S. pneumoniae* polypeptides described in Table 1 as predicted by the inventors. ~~Each~~^{The} *S. pneumoniae* polypeptide shown in Table 1 has one or more antigenic epitopes described in Table 2. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly. The exact location of the antigenic determinant may shift by about 1 to 5 residues, more likely 1 to 2 residues, depending on the criteria used. Thus, the first antigenic determinant described in Table 2, ~~"Lys 1 to Ile-10" of SP001, represents a peptide comprising the lysine at position 1 in SEQ ID NO:2 through and including the isoleucine at position 10 in SEQ ID NO:2, but may include more or fewer residues than those 10.~~ It will also be appreciated that, generally speaking, amino acids can be added to either terminus of a peptide or polypeptide containing an antigenic epitope without affecting its activity, whereas removing residues from a peptide or polypeptide containing only the antigenic determinant is much more likely to destroy activity. It will be appreciated that the residues and locations shown described in Table 2 correspond to the amino acid sequences for ~~each~~^{the} ORF shown in Table 1 and in the Sequence Listing.

Explanation of Table 3

Table 3 shows PCR primers designed by the inventors for the amplification of polynucleotides encoding polypeptides of the present invention according to the method of Example 1. PCR primer design is routine in the art and those shown in Table 3 are provided merely for the convenience of the skilled artisan. It will be appreciated that others can be used with equal success.

For each primer, the table lists the corresponding ORF designation from Table 1 followed by either an "A" or a "B". The "A" primers are the 5' primers and the "B" primers 3'. A restriction enzyme site was built into each primer to allow ease of cloning. The restriction enzyme which will recognize and cleave a sequence within each primer is shown in Table 3, as well, under the heading

"RE" for restriction enzyme. Finally the sequence identifier is shown in Table 3 for each primer for easy correlation with the Sequence Listing.

Selection of Nucleic Acid Sequences Encoding Antigenic S. pneumoniae Polypeptides

The present invention provides a select number of ORFs from those presented in the fragments of the *S. pneumoniae* genome which may prove useful for the generation of a protective immune response. The sequenced *S. pneumoniae* genomic DNA was obtained from a sub-cultured isolate of *S. pneumoniae* Strain 7/87 14.8.91, which has been deposited at the American Type Culture Collection, as a convenience to those of skill in the art. The *S. pneumoniae* isolate was deposited on October 10, 1996 at the ATCC, 12301 ^{10301 University Blvd,} ~~Manassas, VA 20108-2209, USA,~~ ~~Park Lawn Drive, Rockville, Maryland 20852,~~ and given accession number 55840. A genomic library constructed from DNA isolated from the *S. pneumoniae* isolate was also deposited at the ATCC on October 11, 1996 and given ATCC Deposit No. 97755. A more complete listing of the sequence obtained from the *S. pneumoniae* genome may be found in co-pending U.S. Provisional Application Serial No. 60/029,960, filed 10/31/96, incorporated herein by reference in its entirety. Some ORFs contained in the subset of fragments of the *S. pneumoniae* genome disclosed herein were derived through the use of a number of screening criteria detailed below.

The selected ORFs do not consist of complete ORFs. Although a polypeptide representing a complete ORF may be the closest approximation of a protein native to an organism, it is not always preferred to express a complete ORF in a heterologous system. It may be challenging to express and purify a highly hydrophobic protein by common laboratory methods. Thus, the polypeptide vaccine candidates described herein may have been modified slightly to simplify the production of recombinant protein. For example, nucleotide sequences which encode highly hydrophobic domains, such as those found at the amino terminal signal sequence, have been excluded from some constructs used for *in vitro* expression of the polypeptides. Furthermore, any highly hydrophobic amino acid sequences occurring at the carboxy terminus have also been excluded from the recombinant expression constructs. Thus, in one embodiment, a polypeptide which represents a truncated or modified ORF may be used as an antigen.

While numerous methods are known in the art for selecting potentially immunogenic polypeptides, many of the ORFs disclosed herein were selected

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103097

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on the basis of screening all theoretical *S. pneumoniae* ORFs for several aspects of potential immunogenicity. One set of selection criteria are as follows:

1. *Type I signal sequence*: An amino terminal type I signal sequence generally directs a nascent protein across the plasma and outer membranes to the exterior of the bacterial cell. Experimental evidence obtained from studies with *Escherichia coli* suggests that the typical type I signal sequence consists of the following biochemical and physical attributes (Izard, J. W. and Kendall, D. A. *Mol. Microbiol.* **13**:765-773 (1994)). The length of the type I signal sequence is approximately 15 to 25 primarily hydrophobic amino acid residues with a net positive charge in the extreme amino terminus. In addition, the central region of the signal sequence adopts an alpha-helical conformation in a hydrophobic environment. Finally, the region surrounding the actual site of cleavage is ideally six residues long, with small side-chain amino acids in the -1 and -3 positions.

2. *Type IV signal sequence*: The type IV signal sequence is an example of the several types of functional signal sequences which exist in addition to the type I signal sequence detailed above. Although functionally related, the type IV signal sequence possesses a unique set of biochemical and physical attributes (Strom, M. S. and Lory, S., *J. Bacteriol.* **174**:7345-7351 (1992)). These are typically six to eight amino acids with a net basic charge followed by an additional sixteen to thirty primarily hydrophobic residues. The cleavage site of a type IV signal sequence is typically after the initial six to eight amino acids at the extreme amino terminus. In addition, type IV signal sequences generally contain a phenylalanine residue at the +1 site relative to the cleavage site.

3. *Lipoprotein*: Studies of the cleavage sites of twenty-six bacterial lipoprotein precursors has allowed the definition of a consensus amino acid sequence for lipoprotein cleavage. Nearly three-fourths of the bacterial lipoprotein precursors examined contained the sequence L-(A,S)-(G,A)-C at positions -3 to +1, relative to the point of cleavage (Hayashi, S. and Wu, H. C., *J. Bioenerg. Biomembr.* **22**:451-471 (1990)).

4. *LPXTG motif*: It has been experimentally determined that most anchored proteins found on the surface of gram-positive bacteria possess a highly conserved carboxy terminal sequence. More than fifty such proteins from organisms such as *S. pyogenes*, *S. mutans*, *E. faecalis*, *S. pneumoniae*, and others, have been identified based on their extracellular location and carboxy terminal amino acid sequence (Fischetti, V. A., *ASM News* **62**:405-410 (1996)). The conserved region consists of six charged amino acids at the extreme carboxy terminus coupled to 15-20 hydrophobic amino acids

presumed to function as a transmembrane domain. Immediately adjacent to the transmembrane domain is a six amino acid sequence conserved in nearly all proteins examined. The amino acid sequence of this region is L-P-X-T-G-X, where X is any amino acid.

An algorithm for selecting antigenic and immunogenic *S. pneumoniae* polypeptides including the foregoing criteria was developed. Use of the algorithm by the inventors to select immunologically useful *S. pneumoniae* polypeptides resulted in the selection of ~~a number of~~ the disclosed ORFs. Polypeptides comprising the polypeptides identified in this group may be produced by techniques standard in the art and as further described herein.

Nucleic Acid Molecules

The present invention provides isolated nucleic acid molecules comprising polynucleotides encoding the *S. pneumoniae* polypeptides having the amino acid sequences described in Table 1 and shown as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226, which ^{was} ~~were~~ determined by sequencing the genome of *S. pneumoniae* and selected as putative immunogens.

Unless otherwise indicated, all nucleotide sequences determined by sequencing a DNA molecule herein were determined using an automated DNA sequencer (such as the Model 373 from Applied Biosystems, Inc.), and all amino acid sequences of polypeptides encoded by DNA molecules determined herein were predicted by translation of DNA sequences determined as above. Therefore, as is known in the art for any DNA sequence determined by this automated approach, any nucleotide sequence determined herein may contain some errors. Nucleotide sequences determined by automation are typically at least about 90% identical, more typically at least about 95% to at least about 99.9% identical to the actual nucleotide sequence of the sequenced DNA molecule. The actual sequence can be more precisely determined by other approaches including manual DNA sequencing methods well known in the art. As is also known in the art, a single insertion or deletion in a determined nucleotide sequence compared to the actual sequence will cause a frame shift in translation of the nucleotide sequence such that the predicted amino acid sequence encoded by a determined nucleotide sequence will be completely different from the amino acid sequence actually encoded by the sequenced DNA molecule, beginning at the point of such an insertion or deletion.

Unless otherwise indicated, each "nucleotide sequence" set forth herein is presented as a sequence of deoxyribonucleotides (abbreviated A, G, C and

T). However, by "nucleotide sequence" of a nucleic acid molecule or polynucleotide is intended, for a DNA molecule or polynucleotide, a sequence of deoxyribonucleotides, and for an RNA molecule or polynucleotide, the corresponding sequence of ribonucleotides (A, G, C and U), where each thymidine deoxyribonucleotide (T) in the specified deoxyribonucleotide sequence is replaced by the ribonucleotide uridine (U). For instance, reference to an RNA molecule having a sequence described in Table 1 set forth using deoxyribonucleotide abbreviations is intended to indicate an RNA molecule having a sequence in which each deoxyribonucleotide A, G or C described in Table 1 has been replaced by the corresponding ribonucleotide A, G or C, and each deoxyribonucleotide T has been replaced by a ribonucleotide U.

Nucleic acid molecules of the present invention may be in the form of RNA, such as mRNA, or in the form of DNA, including, for instance, cDNA and genomic DNA obtained by cloning or produced synthetically. The DNA may be double-stranded or single-stranded. Single-stranded DNA or RNA may be the coding strand, also known as the sense strand, or it may be the non-coding strand, also referred to as the anti-sense strand.

By "isolated" nucleic acid molecule(s) is intended a nucleic acid molecule, DNA or RNA, which has been removed from its native environment. For example, recombinant DNA molecules contained in a vector are considered isolated for the purposes of the present invention. Further examples of isolated DNA molecules include recombinant DNA molecules maintained in heterologous host cells or purified (partially or substantially) DNA molecules in solution. Isolated RNA molecules include *in vivo* or *in vitro* RNA transcripts of the DNA molecules of the present invention. Isolated nucleic acid molecules according to the present invention further include such molecules produced synthetically.

Isolated nucleic acid molecules of the present invention include DNA molecules comprising a nucleotide sequence described in Table 1 and shown as SEQ ID NO:1, ~~SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225~~, DNA molecules comprising the coding sequences for the polypeptides described in Table 1 and shown as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226; and DNA molecules which comprise sequences substantially different from those described above but which, due to the degeneracy of the genetic code, still encode the *S. pneumoniae* polypeptides described in Table 1. Of course, the genetic code is well known in the art. Thus, it would be routine for one skilled in the art to generate such degenerate variants.

The invention also provides nucleic acid molecules having sequences complementary to ^{that} ~~any one of those~~ described in Table 1. Such isolated molecules, particularly DNA molecules, are useful as probes for detecting expression of *Streptococcal* genes, for instance, by Northern blot analysis or the polymerase chain reaction (PCR).

The present invention is further directed to fragments of the isolated nucleic acid molecules described herein. By a fragment of an isolated nucleic acid molecule having a nucleotide sequence described in Table 1, is intended fragments at least about 15 nt, and more preferably at least about 17 nt, still more preferably at least about 20 nt, and even more preferably, at least about 25 nt in length which are useful as diagnostic probes and primers as discussed herein. Of course, larger fragments 50-100 nt in length are also useful according to the present invention as are fragments corresponding to ~~the~~ ^{most, if} ~~not all, of a~~ nucleotide sequence described in Table 1. By a fragment at least 20 nt in length, for example, is intended fragments which include 20 or more contiguous bases of a nucleotide sequence as described in Table 1. Since the nucleotide sequences identified in Table 1 are provided as SEQ ID NO:1, ~~SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225,~~ ^{SEQ} generating such DNA fragments would be routine to the skilled artisan. For example, such fragments could be generated synthetically.

Preferred nucleic acid fragments of the present invention also include nucleic acid molecules comprising nucleotide sequences encoding epitope-bearing portions of the *S. pneumoniae* polypeptides identified in Table 1. Such nucleic acid fragments of the present invention include, for example, nucleotide sequences encoding polypeptide fragments comprising from about the amino terminal residue to about the carboxy terminal residue of ~~each~~ ^{the} fragment shown in Table 2. The above referred to polypeptide fragments are antigenic regions of the *S. pneumoniae* polypeptides identified in Table 1.

In another aspect, the invention provides isolated nucleic acid molecules comprising polynucleotides which hybridize under stringent hybridization conditions to a portion of a polynucleotide in a nucleic acid molecule of the invention described above, for instance, a nucleic acid sequence identified in Table 1. By "stringent hybridization conditions" is intended overnight incubation at 42°C in a solution comprising: 50% formamide, 5x SSC (150 mM NaCl, 15 mM trisodium citrate), 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and 20 g/ml denatured, sheared salmon sperm DNA, followed by washing the filters in 0.1x SSC at about 65°C.

By polynucleotides which hybridize to a "portion" of a polynucleotide is intended polynucleotides (either DNA or RNA) which hybridize to at least about 15 nucleotides (nt), and more preferably at least about 17 nt, still more preferably at least about 20 nt, and even more preferably about 25-70 nt of the reference polynucleotide. These are useful as diagnostic probes and primers as discussed above and in more detail below.

Of course, polynucleotides hybridizing to a larger portion of the reference polynucleotide, for instance, a portion 50-100 nt in length, or even to the entire length of the reference polynucleotide, are also useful as probes according to the present invention, as are polynucleotides corresponding to ~~most, if not all, of~~ a nucleotide sequence as identified in Table 1. By a portion of a polynucleotide of "at least 20 nt in length," for example, is intended 20 or more contiguous nucleotides from the nucleotide sequence of the reference polynucleotide (e.g., a nucleotide sequence as described in Table 1). As noted above, such portions are useful diagnostically either as probes according to conventional DNA hybridization techniques or as primers for amplification of a target sequence by PCR, as described in the literature (for instance, in *Molecular Cloning, A Laboratory Manual*, 2nd. edition, Sambrook, J., Fritsch, E. F. and Maniatis, T., eds., Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y. (1989), the entire disclosure of which is hereby incorporated herein by reference).

Since nucleic acid sequences encoding the *S. pneumoniae* polypeptides of the present invention are identified in Table 1 and provided as SEQ ID NO:1, ~~SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225,~~ generating polynucleotides which hybridize to portions of these sequences would be routine to the skilled artisan. For example, the hybridizing polynucleotides of the present invention could be generated synthetically according to known techniques.

As indicated, nucleic acid molecules of the present invention which encode *S. pneumoniae* polypeptides of the present invention may include, but are not limited to those encoding the amino acid sequences of the polypeptides by themselves; and additional coding sequences which code for additional amino acids, such as those which provide additional functionalities. Thus, the sequences encoding these polypeptides may be fused to a marker sequence, such as a sequence encoding a peptide which facilitates purification of the fused polypeptide. In certain preferred embodiments of this aspect of the invention, the marker amino acid sequence is a hexa-histidine peptide, such as the tag provided in a pQE vector (Qiagen, Inc.), among others, many of which are

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commercially available. As described by Gentz and colleagues (*Proc. Natl. Acad. Sci. USA* **86**:821-824 (1989)), for instance, hexa-histidine provides for convenient purification of the resulting fusion protein.

Thus, the present invention also includes genetic fusions wherein the *S. pneumoniae* nucleic acid sequences coding sequences identified in Table 1 are linked to additional nucleic acid sequences to produce fusion proteins. These fusion proteins may include epitopes of streptococcal or non-streptococcal origin designed to produce proteins having enhanced immunogenicity. Further, the fusion proteins of the present invention may contain antigenic determinants known to provide helper T-cell stimulation, peptides encoding sites for post-translational modifications which enhance immunogenicity (*e.g.*, acylation), peptides which facilitate purification (*e.g.*, histidine "tag"), or amino acid sequences which target the fusion protein to a desired location (*e.g.*, a heterologous leader sequence).

In all cases of bacterial expression, an N-terminal methionine residues is added. In many cases, however, the N-terminal methionine residues is cleaved off post-translationally. Thus, the invention includes polypeptides shown in Table 1 with, and without an N-terminal methionine.

The present invention thus includes nucleic acid molecules and sequences which encode fusion proteins comprising one or more *S. pneumoniae* polypeptides of the present invention fused to an amino acid sequence which allows for post-translational modification to enhance immunogenicity. This post-translational modification may occur either *in vitro* or when the fusion protein is expressed *in vivo* in a host cell. An example of such a modification is the introduction of an amino acid sequence which results in the attachment of a lipid moiety.

Thus, as indicated above, the present invention includes genetic fusions wherein a *S. pneumoniae* nucleic acid sequence identified in Table 1 is linked to a nucleotide sequence encoding another amino acid sequence. These other amino acid sequences may be of streptococcal origin (*e.g.*, another sequence selected from Table 1) or non-streptococcal origin.

The present invention further relates to variants of the nucleic acid molecules of the present invention, which encode portions, analogs or derivatives of the *S. pneumoniae* polypeptides described in Table 1. Variants may occur naturally, such as a natural allelic variant. By an "allelic variant" is intended one of several alternate forms of a gene occupying a given locus on a chromosome of an organism (*Genes II*, Lewin, B., ed., John Wiley & Sons,

New York (1985)). Non-naturally occurring variants may be produced using art-known mutagenesis techniques.

Such variants include those produced by nucleotide substitutions, deletions or additions. The substitutions, deletions or additions may involve one or more nucleotides. These variants may be altered in coding regions, non-coding regions, or both. Alterations in the coding regions may produce conservative or non-conservative amino acid substitutions, deletions or additions. Especially preferred among these are silent substitutions, additions and deletions, which do not alter the properties and activities of the *S. pneumoniae* polypeptides disclosed herein or portions thereof. Silent ~~substitutions~~ ^{substitutions} are most likely to be made in non-epitopic regions. Guidance regarding those regions containing epitopes is provided herein, for example, in Table 2. Also especially preferred in this regard are conservative substitutions.

Further embodiments of the invention include isolated nucleic acid molecules comprising a polynucleotide having a nucleotide sequence at least 90% identical, and more preferably at least 95%, 96%, 97%, 98% or 99% identical to: (a) a nucleotide sequence encoding ~~any of~~ the amino acid sequences of the polypeptides identified in Table 1; and (b) a nucleotide sequence complementary to ~~any of~~ the nucleotide sequences in (a) above.

By a polynucleotide having a nucleotide sequence at least, for example, 95% "identical" to a reference nucleotide sequence encoding a *S. pneumoniae* polypeptide described in Table 1, is intended that the nucleotide sequence of the polynucleotide is identical to the reference sequence except that the polynucleotide sequence may include up to five point mutations per each 100 nucleotides of the reference nucleotide sequence encoding the subject *S. pneumoniae* polypeptide. In other words, to obtain a polynucleotide having a nucleotide sequence at least 95% identical to a reference nucleotide sequence, up to 5% of the nucleotides in the reference sequence may be deleted or substituted with another nucleotide, or a number of nucleotides up to 5% of the total nucleotides in the reference sequence may be inserted into the reference sequence. These mutations of the reference sequence may occur at the 5' or 3' terminal positions of the reference nucleotide sequence or anywhere between those terminal positions, interspersed either individually among nucleotides in the reference sequence or in one or more contiguous groups within the reference sequence.

Certain nucleotides within ~~some of the nucleic acid sequences~~ ^{the nucleic acid sequence} shown in Table 1 were ambiguous upon sequencing. Completely unknown sequences are shown as an "N". Other unresolved nucleotides are known to be either a

purine, shown as "R", or a pyrimidine, shown as "Y". Accordingly, when determining identity between two nucleotide sequences, identity is met where any nucleotide, including an "R", "Y" or "N", is found in a test sequence and at the corresponding position in the ^{reference} ~~reference~~ sequence (from Table 1). Likewise, an A, G or "R" in a test sequence is identical to an "R" in the reference sequence; and a T, C or "Y" in a test sequence is identical to a "Y" in the reference sequence.

As a practical matter, whether any particular nucleic acid molecule is at least 90%, 95%, 96%, 97%, 98% or 99% identical to, for instance, a nucleotide sequence described in Table 1 can be determined conventionally using known computer programs such as the Bestfit program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, WI 53711). Bestfit uses the local homology algorithm of Smith and Waterman (*Advances in Applied Mathematics* 2:482-489 (1981)), to find the best segment of homology between two sequences. When using Bestfit or any other sequence alignment program to determine whether a particular sequence is, for instance, 95% identical to a reference sequence according to the present invention, the parameters are set, of course, such that the percentage of identity is calculated over the full length of the reference nucleotide sequence and that gaps in homology of up to 5% of the total number of nucleotides in the reference sequence are allowed.

The present application is directed to nucleic acid molecules at least 90%, 95%, 96%, 97%, 98% or 99% identical to a nucleic acid sequence described in Table 1. One of skill in the art would still know how to use the nucleic acid molecule, for instance, as a hybridization probe or a polymerase chain reaction (PCR) primer. Uses of the nucleic acid molecules of the present invention include, *inter alia*, (1) isolating *Streptococcal* genes or allelic variants thereof from either a genomic or cDNA library and (2) Northern Blot or PCR analysis for detecting *Streptococcal* mRNA expression.

Of course, due to the degeneracy of the genetic code, one of ordinary skill in the art will immediately recognize that a large number of nucleic acid molecules having a sequence at least 90%, 95%, 96%, 97%, 98%, or 99% identical to a nucleic acid sequence identified in Table 1 will encode the same polypeptide. In fact, since degenerate variants of these nucleotide sequences all encode the same polypeptide, this will be clear to the skilled artisan even without performing the above described comparison assay.

It will be further recognized in the art that, for such nucleic acid molecules that are not degenerate variants, a reasonable number will also encode

proteins having antigenic epitopes of the *S. pneumoniae* polypeptides of the present invention. This is because the skilled artisan is fully aware of amino acid substitutions that are either less likely or not likely to significantly effect the antigenicity of a polypeptide (*e.g.*, replacement of an amino acid in a region which is not believed to form an antigenic epitope). For example, since antigenic epitopes have been identified which contain as few as six amino acids (see Harlow, *et al.*, *Antibodies: A Laboratory Manual*, 2nd Ed.; Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York (1988), page 76), in instances where a polypeptide has multiple antigenic epitopes the alteration of several amino acid residues would often not be expected to eliminate all of the antigenic epitopes of that polypeptide. This is especially so when the alterations are in regions believed to not constitute antigenic epitopes.

Vectors and Host Cells

The present invention also relates to vectors which include the isolated DNA molecules of the present invention, host cells which are genetically engineered with the recombinant vectors, and the production of *S. pneumoniae* polypeptides or fragments thereof by recombinant techniques.

Recombinant constructs may be introduced into host cells using well known techniques such as infection, transduction, transfection, transvection, electroporation and transformation. The vector may be, for example, a phage, plasmid, viral or retroviral vector. Retroviral vectors may be replication competent or replication defective. In the latter case, viral propagation generally will occur only in complementing host cells.

The polynucleotides may be joined to a vector containing a selectable marker for propagation in a host. Generally, a plasmid vector is introduced in a precipitate, such as a calcium phosphate precipitate, or in a complex with a charged lipid. If the vector is a virus, it may be packaged *in vitro* using an appropriate packaging cell line and then transduced into host cells.

Preferred are vectors comprising *cis*-acting control regions to the polynucleotide of interest. Appropriate *trans*-acting factors may be supplied by the host, supplied by a complementing vector or supplied by the vector itself upon introduction into the host.

In certain preferred embodiments in this regard, the vectors provide for specific expression, which may be inducible and/or cell type-specific. Particularly preferred among such vectors are those inducible by environmental factors that are easy to manipulate, such as temperature and nutrient additives.

Expression vectors useful in the present invention include chromosomal-, episomal- and virus-derived vectors, *e.g.*, vectors derived from bacterial plasmids, bacteriophage, yeast episomes, yeast chromosomal elements, viruses such as baculoviruses, papova viruses, vaccinia viruses, adenoviruses, fowl pox viruses, pseudorabies viruses and retroviruses, and vectors derived from combinations thereof, such as cosmids and phagemids.

The DNA insert should be operatively linked to an appropriate promoter, such as the phage lambda PL promoter, the *E. coli lac*, *trp* and *tac* promoters, the SV40 early and late promoters and promoters of retroviral LTRs, to name a few. Other suitable promoters will be known to the skilled artisan. The expression constructs will further contain sites for transcription initiation, termination and, in the transcribed region, a ribosome binding site for translation. The coding portion of the mature transcripts expressed by the constructs will preferably include a translation initiating site at the beginning and a termination codon (UAA, UGA or UAG) appropriately positioned at the end of the polypeptide to be translated.

As indicated, the expression vectors will preferably include at least one selectable marker. Such markers include dihydrofolate reductase or neomycin resistance for eukaryotic cell culture and tetracycline or ampicillin resistance genes for culturing in *E. coli* and other bacteria. Representative examples of appropriate hosts include, but are not limited to, bacterial cells, such as *E. coli*, *Streptomyces* and *Salmonella typhimurium* cells; fungal cells, such as yeast cells; insect cells such as *Drosophila* S2 and *Spodoptera* Sf9 cells; animal cells such as CHO, COS and Bowes melanoma cells; and plant cells. Appropriate culture mediums and conditions for the above-described host cells are known in the art.

Among vectors preferred for use in bacteria include pQE70, pQE60 and pQE-9, available from Qiagen; pBS vectors, Phagescript vectors, Bluescript vectors, pNH8A, pNH16a, pNH18A, pNH46A available from Stratagene; pET series of vectors available from Novagen; and ptrc99a, pKK223-3, pKK233-3, pDR540, pRIT5 available from Pharmacia. Among preferred eukaryotic vectors are pWLNEO, pSV2CAT, pOG44, pXT1 and pSG available from Stratagene; and pSVK3, pBPV, pMSG and pSVL available from Pharmacia. Other suitable vectors will be readily apparent to the skilled artisan.

Among known bacterial promoters suitable for use in the present invention include the *E. coli lacI* and *lacZ* promoters, the T3 and T7 promoters, the *gpt* promoter, the lambda PR and PL promoters and the *trp* promoter. Suitable eukaryotic promoters include the CMV immediate early promoter, the

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HSV thymidine kinase promoter, the early and late SV40 promoters, the promoters of retroviral LTRs, such as those of the Rous sarcoma virus (RSV), and metallothionein promoters, such as the mouse metallothionein-I promoter.

Introduction of the construct into the host cell can be effected by calcium phosphate transfection, DEAE-dextran mediated transfection, cationic lipid-mediated transfection, electroporation, transduction, infection or other methods. Such methods are described in many standard laboratory manuals (for example, Davis, *et al.*, *Basic Methods In Molecular Biology* (1986)).

Transcription of DNA encoding the polypeptides of the present invention by higher eukaryotes may be increased by inserting an enhancer sequence into the vector. Enhancers are *cis*-acting elements of DNA, usually about from 10 to 300 bp that act to increase transcriptional activity of a promoter in a given host cell-type. Examples of enhancers include the SV40 enhancer, which is located on the late side of the replication origin at bp 100 to 270, the cytomegalovirus early promoter enhancer, the polyoma enhancer on the late side of the replication origin, and adenovirus enhancers.

For secretion of the translated polypeptide into the lumen of the endoplasmic reticulum, into the periplasmic space or into the extracellular environment, appropriate secretion signals may be incorporated into the expressed polypeptide. The signals may be endogenous to the polypeptide or they may be heterologous signals.

The polypeptide may be expressed in a modified form, such as a fusion protein, and may include not only secretion signals, but also additional heterologous functional regions. For instance, a region of additional amino acids, particularly charged amino acids, may be added to the N-terminus of the polypeptide to improve stability and persistence in the host cell, during purification, or during subsequent handling and storage. Also, peptide moieties may be added to the polypeptide to facilitate purification. Such regions may be removed prior to final preparation of the polypeptide. The addition of peptide moieties to polypeptides to engender secretion or excretion, to improve stability and to facilitate purification, among others, are familiar and routine techniques in the art. A preferred fusion protein comprises a heterologous region from immunoglobulin that is useful to solubilize proteins. For example, EP-A-O 464 533 (Canadian counterpart 2045869) discloses fusion proteins comprising various portions of constant region of immunoglobulin molecules together with another human protein or part thereof. In many cases, the Fc part in a fusion protein is thoroughly advantageous for use in therapy and diagnosis and thus results, for example, in improved pharmacokinetic properties (EP-A 0232 262).

On the other hand, for some uses it would be desirable to be able to delete the Fc part after the fusion protein has been expressed, detected and purified in the advantageous manner described. This is the case when Fc portion proves to be a hindrance to use in therapy and diagnosis, for example when the fusion protein is to be used as antigen for immunizations. In drug discovery, for example, human proteins, such as, hIL5-receptor has been fused with Fc portions for the purpose of high-throughput screening assays to identify antagonists of hIL-5. See Bennett, D. *et al.*, *J. Molec. Recogn.* 8:52-58 (1995) and Johanson, K. *et al.*, *J. Biol. Chem.* 270 (16):9459-9471 (1995).

The *S. pneumoniae* polypeptides can be recovered and purified from recombinant cell cultures by well-known methods including ammonium sulfate or ethanol precipitation, acid extraction, anion or cation exchange chromatography, phosphocellulose chromatography, hydrophobic interaction chromatography, affinity chromatography, hydroxylapatite chromatography, lectin chromatography and high performance liquid chromatography ("HPLC") is employed for purification. Polypeptides of the present invention include naturally purified products, products of chemical synthetic procedures, and products produced by recombinant techniques from a prokaryotic or eukaryotic host, including, for example, bacterial, yeast, higher plant, insect and mammalian cells.

Polypeptides and Fragments

The invention further provides isolated polypeptides having the amino acid sequences described in Table 1, and shown as SEQ ID NO:2, ~~SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226,~~ and peptides or polypeptides comprising portions of the above polypeptides. The terms "peptide" and "oligopeptide" are considered synonymous (as is commonly recognized) and each term can be used interchangeably as the context requires to indicate a chain of at least two amino acids coupled by peptidyl linkages. The word "polypeptide" is used herein for chains containing more than ten amino acid residues. All oligopeptide and polypeptide formulas or sequences herein are written from left to right and in the direction from amino terminus to carboxy terminus.

Some amino acid sequences of the *S. pneumoniae* polypeptides described in Table 1 can be varied without significantly effecting the antigenicity of the polypeptides. If such differences in sequence are contemplated, it should be remembered that there will be critical areas on the polypeptide which determine antigenicity. In general, it is possible to replace residues which do

not form part of an antigenic epitope without significantly effecting the antigenicity of a polypeptide. Guidance for such alterations is given in Table 2 wherein epitopes for ~~each~~^{the} polypeptide ~~is~~^{are} delineated.

The polypeptides of the present invention are preferably provided in an isolated form. By "isolated polypeptide" is intended a polypeptide removed from its native environment. Thus, a polypeptide produced and/or contained within a recombinant host cell is considered isolated for purposes of the present invention. Also intended as an "isolated polypeptide" is a polypeptide that has been purified, partially or substantially, from a recombinant host cell. For example, recombinantly produced versions of the *S. pneumoniae* polypeptides described in Table 1 can be substantially purified by the one-step method described by Smith and Johnson (*Gene* 67:31-40 (1988)).

The polypeptides of the present invention include: (a) an amino acid sequence of ~~any of~~ the polypeptides described in Table 1; and (b) an amino acid sequence of an epitope-bearing portion of ~~any one of~~ the polypeptides of (a); as well as polypeptides with at least 70% similarity, and more preferably at least 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% similarity to those described in (a) or (b) above, as well as polypeptides having an amino acid sequence at least 70% identical, more preferably at least 75% identical, and still more preferably 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% identical to those above.

By "% similarity" for two polypeptides is intended a similarity score produced by comparing the amino acid sequences of the two polypeptides using the Bestfit program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, WI 53711) and the default settings for determining similarity. Bestfit uses the local homology algorithm of Smith and Waterman (*Advances in Applied Mathematics* 2:482-489 (1981)) to find the best segment of similarity between two sequences.

By a polypeptide having an amino acid sequence at least, for example, 95% "identical" to a reference amino acid sequence of a *S. pneumoniae* polypeptide is intended that the amino acid sequence of the polypeptide is identical to the reference sequence except that the polypeptide sequence may include up to five amino acid alterations per each 100 amino acids of the reference amino acid sequence. In other words, to obtain a polypeptide having an amino acid sequence at least 95% identical to a reference amino acid sequence, up to 5% of the amino acid residues in the reference sequence may be deleted or substituted with another amino acid, or a number of amino acids up to

5% of the total amino acid residues in the reference sequence may be inserted into the reference sequence. These alterations of the reference sequence may occur at the amino or carboxy terminal positions of the reference amino acid sequence or anywhere between those terminal positions, interspersed either individually among residues in the reference sequence or in one or more contiguous groups within the reference sequence.

The amino acid sequences shown in Table 1 may have ^{one} or more "X" residues. "X" represents unknown. Thus, for purposes of defining identity, if any amino acid is present at the same position in a reference amino acid sequence (shown in Table 1) where an X is shown, the two sequences are identical at that position.

As a practical matter, whether any particular polypeptide is at least 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% identical to, for instance, an amino acid sequence shown in Table 1, can be determined conventionally using known computer programs such the Bestfit⁹ program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, WI 53711). When using Bestfit or any other sequence alignment program to determine whether a particular sequence is, for instance, 95% identical to a reference sequence according to the present invention, the parameters are set, of course, such that the percentage of identity is calculated over the full length of the reference amino acid sequence and that gaps in homology of up to 5% of the total number of amino acid residues in the reference sequence are allowed.

As described below, the polypeptides of the present invention can also be used to raise polyclonal and monoclonal antibodies, which are useful in assays for detecting *Streptococcal* protein expression.

In another aspect, the invention provides peptides and polypeptides comprising epitope-bearing portions of the *S. pneumoniae* polypeptides of the invention. These epitopes are immunogenic or antigenic epitopes of the polypeptides of the invention. An "immunogenic epitope" is defined as a part of a protein that elicits an antibody response when the whole protein or polypeptide is the immunogen. These immunogenic epitopes are believed to be confined to a few loci on the molecule. On the other hand, a region of a protein molecule to which an antibody can bind is defined as an "antigenic determinant" or "antigenic epitope." The number of immunogenic epitopes of a protein generally is less than the number of antigenic epitopes (Geysen, *et al.*, *Proc. Natl. Acad. Sci. USA* **81**:3998- 4002 (1983)). Predicted antigenic epitopes are shown in Table 2, below.

As to the selection of peptides or polypeptides bearing an antigenic epitope (*i.e.*, that contain a region of a protein molecule to which an antibody can bind), it is well known in that art that relatively short synthetic peptides that mimic part of a protein sequence are routinely capable of eliciting an antiserum that reacts with the partially mimicked protein (for instance, Sutcliffe, J., *et al.*, *Science* **219**:660-666 (1983)). Peptides capable of eliciting protein-reactive sera are frequently represented in the primary sequence of a protein, can be characterized by a set of simple chemical rules, and are confined neither to immunodominant regions of intact proteins (*i.e.*, immunogenic epitopes) nor to the amino or carboxyl terminals. Peptides that are extremely hydrophobic and those of six or fewer residues generally are ineffective at inducing antibodies that bind to the mimicked protein; longer, peptides, especially those containing proline residues, usually are effective (Sutcliffe, *et al.*, *supra*, p. 661). For instance, 18 of 20 peptides designed according to these guidelines, containing 8-39 residues covering 75% of the sequence of the influenza virus hemagglutinin HA1 polypeptide chain, induced antibodies that reacted with the HA1 protein or intact virus; and 12/12 peptides from the MuLV polymerase and 18/18 from the rabies glycoprotein induced antibodies that precipitated the respective proteins.

Antigenic epitope-bearing peptides and polypeptides of the invention are therefore useful to raise antibodies, including monoclonal antibodies, that bind specifically to a polypeptide of the invention. Thus, a high proportion of hybridomas obtained by fusion of spleen cells from donors immunized with an antigen epitope-bearing peptide generally secrete antibody reactive with the native protein (Sutcliffe, *et al.*, *supra*, p. 663). The antibodies raised by antigenic epitope-bearing peptides or polypeptides are useful to detect the mimicked protein, and antibodies to different peptides may be used for tracking the fate of various regions of a protein precursor which undergoes post-translational processing. The peptides and anti-peptide antibodies may be used in a variety of qualitative or quantitative assays for the mimicked protein, for instance in competition assays since it has been shown that even short peptides (*e.g.*, about 9 amino acids) can bind and displace the larger peptides in immunoprecipitation assays (for instance, Wilson, *et al.*, *Cell* **37**:767-778 (1984) p. 777). The anti-peptide antibodies of the invention also are useful for purification of the mimicked protein, for instance, by adsorption chromatography using methods well known in the art.

Antigenic epitope-bearing peptides and polypeptides of the invention designed according to the above guidelines preferably contain a sequence of at

least seven, more preferably at least nine and most preferably between about 15 to about 30 amino acids contained within the amino acid sequence of a polypeptide of the invention. However, peptides or polypeptides comprising a larger portion of an amino acid sequence of a polypeptide of the invention, containing about 30 to about 50 amino acids, or any length up to and including the entire amino acid sequence of a polypeptide of the invention, also are considered epitope-bearing peptides or polypeptides of the invention and also are useful for inducing antibodies that react with the mimicked protein. Preferably, the amino acid sequence of the epitope-bearing peptide is selected to provide substantial solubility in aqueous solvents (*i.e.*, the sequence includes relatively hydrophilic residues and highly hydrophobic sequences are preferably avoided); and sequences containing proline residues are particularly preferred.

Non-limiting examples of antigenic polypeptides or peptides that can be used to generate *Streptococcal*-specific antibodies include portions of the amino acid sequence identified in Table 1. More specifically, Table 2 discloses antigenic fragments of polypeptides of the present invention, which antigenic fragments comprise amino acid sequences from about the first amino acid residues indicated to about the last amino acid residue indicated for each fragment. The polypeptide fragments disclosed in Table 2 are believed to be antigenic regions of the *S. pneumoniae* polypeptide described in Table 1. Thus the invention further includes isolated peptides and polypeptides comprising an amino acid sequence of an epitope shown in Table 2 and polynucleotides encoding said polypeptides.

The epitope-bearing peptides and polypeptides of the invention may be produced by any conventional means for making peptides or polypeptides including recombinant means using nucleic acid molecules of the invention. For instance, an epitope-bearing amino acid sequence of the present invention may be fused to a larger polypeptide which acts as a carrier during recombinant production and purification, as well as during immunization to produce anti-peptide antibodies. Epitope-bearing peptides also may be synthesized using known methods of chemical synthesis. For instance, Houghten has described a simple method for synthesis of large numbers of peptides, such as 10-20 mg of 248 different 13 residue peptides representing single amino acid variants of a segment of the HA1 polypeptide which were prepared and characterized (by ELISA-type binding studies) in less than four weeks (Houghten, R. A. Proc. Natl. Acad. Sci. USA **82**:5131-5135 (1985)). This "Simultaneous Multiple Peptide Synthesis (SMPS)" process is further described in U.S. Patent No. 4,631,211 to Houghten and coworkers (1986). In this procedure the individual

resins for the solid-phase synthesis of various peptides are contained in separate solvent-permeable packets, enabling the optimal use of the many identical repetitive steps involved in solid-phase methods. A completely manual procedure allows 500-1000 or more syntheses to be conducted simultaneously (Houghten, *et al.*, *supra*, p. 5134).

Epitope-bearing peptides and polypeptides of the invention are used to induce antibodies according to methods well known in the art (for instance, Sutcliffe, *et al.*, *supra*; Wilson, *et al.*, *supra*; Chow, M., *et al.*, *Proc. Natl. Acad. Sci. USA* **82**:910-914; and Bittle, F. J., *et al.*, *J. Gen. Virol.* **66**:2347-2354 (1985)). Generally, animals may be immunized with free peptide; however, anti-peptide antibody titer may be boosted by coupling of the peptide to a macromolecular carrier, such as keyhole limpet hemacyanin (KLH) or tetanus toxoid. For instance, peptides containing cysteine may be coupled to carrier using a linker such as m-maleimidobenzoyl-N-hydroxysuccinimide ester (MBS), while other peptides may be coupled to carrier using a more general linking agent such as glutaraldehyde. Animals such as rabbits, rats and mice are immunized with either free or carrier-coupled peptides, for instance, by intraperitoneal and/or intradermal injection of emulsions containing about 100 µg peptide or carrier protein and Freund's adjuvant. Several booster injections may be needed, for instance, at intervals of about two weeks, to provide a useful titer of anti-peptide antibody which can be detected, for example, by ELISA assay using free peptide adsorbed to a solid surface. The titer of anti-peptide antibodies in serum from an immunized animal may be increased by selection of anti-peptide antibodies, for instance, by adsorption to the peptide on a solid support and elution of the selected antibodies according to methods well known in the art.

Immunogenic epitope-bearing peptides of the invention, *i.e.*, those parts of a protein that elicit an antibody response when the whole protein is the immunogen, are identified according to methods known in the art. For instance, Geysen, *et al.*, *supra*, discloses a procedure for rapid concurrent synthesis on solid supports of hundreds of peptides of sufficient purity to react in an enzyme-linked immunosorbent assay. Interaction of synthesized peptides with antibodies is then easily detected without removing them from the support. In this manner a peptide bearing an immunogenic epitope of a desired protein may be identified routinely by one of ordinary skill in the art. For instance, the immunologically important epitope in the coat protein of foot-and-mouth disease virus was located by Geysen *et al. supra* with a resolution of seven amino acids by synthesis of an overlapping set of all 208 possible hexapeptides covering the

entire 213 amino acid sequence of the protein. Then, a complete replacement set of peptides in which all 20 amino acids were substituted in turn at every position within the epitope were synthesized, and the particular amino acids conferring specificity for the reaction with antibody were determined. Thus, peptide
5 analogs of the epitope-bearing peptides of the invention can be made routinely by this method. U.S. Patent No. 4,708,781 to Geysen (1987) further describes this method of identifying a peptide bearing an immunogenic epitope of a desired protein.

Further still, U.S. Patent No. 5,194,392, to Geysen (1990), describes a
10 general method of detecting or determining the sequence of monomers (amino acids or other compounds) which is a topological equivalent of the epitope (*i.e.*, a "mimotope") which is complementary to a particular paratope (antigen binding site) of an antibody of interest. More generally, U.S. Patent No. 4,433,092, also to Geysen (1989), describes a method of detecting or determining a
15 sequence of monomers which is a topographical equivalent of a ligand which is complementary to the ligand binding site of a particular receptor of interest. Similarly, U.S. Patent No. 5,480,971 to Houghten, R. A. *et al.* (1996) discloses linear C₁-C₇-alkyl peralkylated oligopeptides and sets and libraries of such peptides, as well as methods for using such oligopeptide sets and libraries
20 for determining the sequence of a peralkylated oligopeptide that preferentially binds to an acceptor molecule of interest. Thus, non-peptide analogs of the epitope-bearing peptides of the invention also can be made routinely by these methods.

The entire disclosure of each document cited in this section on
25 "Polypeptides and Fragments" is hereby incorporated herein by reference.

As one of skill in the art will appreciate, the polypeptides of the present invention and the epitope-bearing fragments thereof described above can be combined with parts of the constant domain of immunoglobulins (IgG), resulting in chimeric polypeptides. These fusion proteins facilitate purification
30 and show an increased half-life *in vivo*. This has been shown, *e.g.*, for chimeric proteins consisting of the first two domains of the human CD4-polypeptide and various domains of the constant regions of the heavy or light chains of mammalian immunoglobulins (EPA 0,394,827; Traunecker *et al.*, *Nature* 331:84-86 (1988)). Fusion proteins that have a disulfide-linked
35 dimeric structure due to the IgG part can also be more efficient in binding and neutralizing other molecules than a monomeric *S. pneumoniae* polypeptide or

fragment thereof alone (Fountoulakis *et al.*, *J. Biochem.* 270:3958-3964 (1995)).

Diagnostic Assays

The present invention further relates to a method for assaying for *Streptococcal* infection in an animal *via* detecting the expression of genes encoding *Streptococcal* polypeptides (*e.g.*, the polypeptides described Table 1). This method comprises analyzing tissue or body fluid from the animal for *Streptococcus*-specific antibodies or *Streptococcal* nucleic acids or proteins. Analysis of nucleic acid specific to *Streptococcus* can be done by PCR or hybridization techniques using nucleic acid sequences of the present invention as either hybridization probes or primers (*cf. Molecular Cloning: A Laboratory Manual, second edition*, edited by Sambrook, Fritsch, & Maniatis, Cold Spring Harbor Laboratory, 1989; Ereemeeva *et al.*, *J. Clin. Microbiol.* 32:803-810 (1994) which describes differentiation among spotted fever group *Rickettsiae* species by analysis of restriction fragment length polymorphism of PCR-amplified DNA). Methods for detecting *B. burgdorferi* nucleic acids *via* PCR are described, for example, in Chen *et al.*, *J. Clin. Microbiol.* 32:589-595 (1994).

Where diagnosis of a disease state related to infection with *Streptococcus* has already been made, the present invention is useful for monitoring progression or regression of the disease state whereby patients exhibiting enhanced *Streptococcus* gene expression will experience a worse clinical outcome relative to patients expressing these gene(s) at a lower level.

By "assaying for *Streptococcal* infection in an animal *via* detection of genes encoding *Streptococcal* polypeptides" is intended qualitatively or quantitatively measuring or estimating the level of one or more *Streptococcus* polypeptides or the level of nucleic acid encoding *Streptococcus* polypeptides in a first biological sample either directly (*e.g.*, by determining or estimating absolute protein level or nucleic level) or relatively (*e.g.*, by comparing to the *Streptococcus* polypeptide level or mRNA level in a second biological sample). The *Streptococcus* polypeptide level or nucleic acid level in the second sample used for a relative comparison may be undetectable if obtained from an animal which is not infected with *Streptococcus*. When monitoring the progression or regression of a disease state, the *Streptococcus* polypeptide level or nucleic acid level may be compared to a second sample obtained from either an animal infected with *Streptococcus* or the same animal from which the first sample was obtained but taken from that animal at a different time than the first. As will be

appreciated in the art, once a standard *Streptococcus* polypeptide level or nucleic acid level which corresponds to a particular stage of a *Streptococcus* infection is known, it can be used repeatedly as a standard for comparison.

By "biological sample" is intended any biological sample obtained from an animal, cell line, tissue culture, or other source which contains *Streptococcus* polypeptide, mRNA, or DNA. Biological samples include body fluids (such as plasma and synovial fluid) which contain *Streptococcus* polypeptides, and muscle, skin, and cartilage tissues. Methods for obtaining tissue biopsies and body fluids are well known in the art.

The present invention is useful for detecting diseases related to *Streptococcus* infections in animals. Preferred animals include monkeys, apes, cats, dogs, cows, pigs, mice, horses, rabbits and humans. Particularly preferred are humans.

Total RNA can be isolated from a biological sample using any suitable technique such as the single-step guanidinium-thiocyanate-phenol-chloroform method described in Chomczynski and Sacchi, *Anal. Biochem.* 162:156-159 (1987). mRNA encoding *Streptococcus* polypeptides having sufficient homology to the nucleic acid sequences identified in Table 1 to allow for hybridization between complementary sequences are then assayed using any appropriate method. These include Northern blot analysis, S1 nuclease mapping, the polymerase chain reaction (PCR), reverse transcription in combination with the polymerase chain reaction (RT-PCR), and reverse transcription in combination with the ligase chain reaction (RT-LCR).

Northern blot analysis can be performed as described in Harada *et al.*, *Cell* 63:303-312 (1990). Briefly, total RNA is prepared from a biological sample as described above. For the Northern blot, the RNA is denatured in an appropriate buffer (such as glyoxal/dimethyl sulfoxide/sodium phosphate buffer), subjected to agarose gel electrophoresis, and transferred onto a nitrocellulose filter. After the RNAs have been linked to the filter by a UV linker, the filter is prehybridized in a solution containing formamide, SSC, Denhardt's solution, denatured salmon sperm, SDS, and sodium phosphate buffer. A *S. pneumoniae* polypeptide DNA sequence shown in Table 1 labeled according to any appropriate method (such as the ³²P-multiprimered DNA labeling system (Amersham)) is used as probe. After hybridization overnight, the filter is washed and exposed to x-ray film. DNA for use as probe according to the present invention is described in the sections above and will preferably at least 15 bp in length.

S1 mapping can be performed as described in Fujita *et al.*, *Cell* 49:357-367 (1987). To prepare probe DNA for use in S1 mapping, the sense strand of an above-described *S. pneumoniae* DNA sequence of the present invention is used as a template to synthesize labeled antisense DNA. The antisense DNA can then be digested using an appropriate restriction endonuclease to generate further DNA probes of a desired length. Such antisense probes are useful for visualizing protected bands corresponding to the target mRNA (*i.e.*, mRNA encoding *Streptococcus* polypeptides).

Preferably, levels of mRNA encoding *Streptococcus* polypeptides are assayed using the RT-PCR method described in Makino *et al.*, *Technique* 2:295-301 (1990). By this method, the radioactivities of the "amplicons" in the polyacrylamide gel bands are linearly related to the initial concentration of the target mRNA. Briefly, this method involves adding total RNA isolated from a biological sample in a reaction mixture containing a RT primer and appropriate buffer. After incubating for primer annealing, the mixture can be supplemented with a RT buffer, dNTPs, DTT, RNase inhibitor and reverse transcriptase. After incubation to achieve reverse transcription of the RNA, the RT products are then subject to PCR using labeled primers. Alternatively, rather than labeling the primers, a labeled dNTP can be included in the PCR reaction mixture. PCR amplification can be performed in a DNA thermal cycler according to conventional techniques. After a suitable number of rounds to achieve amplification, the PCR reaction mixture is electrophoresed on a polyacrylamide gel. After drying the gel, the radioactivity of the appropriate bands (corresponding to the mRNA encoding the *Streptococcus* polypeptides)) is quantified using an imaging analyzer. RT and PCR reaction ingredients and conditions, reagent and gel concentrations, and labeling methods are well known in the art. Variations on the RT-PCR method will be apparent to the skilled artisan.

Assaying *Streptococcus* polypeptide levels in a biological sample can occur using any art-known method. Preferred for assaying *Streptococcus* polypeptide levels in a biological sample are antibody-based techniques. For example, *Streptococcus* polypeptide expression in tissues can be studied with classical immunohistological methods. In these, the specific recognition is provided by the primary antibody (polyclonal or monoclonal) but the secondary detection system can utilize fluorescent, enzyme, or other conjugated secondary antibodies. As a result, an immunohistological staining of tissue section for pathological examination is obtained. Tissues can also be extracted, *e.g.*, with urea and neutral detergent, for the liberation of *Streptococcus* polypeptides for

Western-blot or dot/slot assay (Jalkanen, M., *et al.*, *J. Cell. Biol.* 101:976-985 (1985); Jalkanen, M., *et al.*, *J. Cell. Biol.* 105:3087-3096 (1987)). In this technique, which is based on the use of cationic solid phases, quantitation of a *Streptococcus* polypeptide can be accomplished using an isolated *Streptococcus* polypeptide as a standard. This technique can also be applied to body fluids.

Other antibody-based methods useful for detecting *Streptococcus* polypeptide gene expression include immunoassays, such as the enzyme linked immunosorbent assay (ELISA) and the radioimmunoassay (RIA). For example, a *Streptococcus* polypeptide-specific monoclonal antibodies can be used both as an immunoabsorbent and as an enzyme-labeled probe to detect and quantify a *Streptococcus* polypeptide. The amount of a *Streptococcus* polypeptide present in the sample can be calculated by reference to the amount present in a standard preparation using a linear regression computer algorithm. Such an ELISA for detecting a tumor antigen is described in Iacobelli *et al.*, *Breast Cancer Research and Treatment* 11:19-30 (1988). In another ELISA assay, two distinct specific monoclonal antibodies can be used to detect *Streptococcus* polypeptides in a body fluid. In this assay, one of the antibodies is used as the immunoabsorbent and the other as the enzyme-labeled probe.

The above techniques may be conducted essentially as a "one-step" or "two-step" assay. The "one-step" assay involves contacting the *Streptococcus* polypeptide with immobilized antibody and, without washing, contacting the mixture with the labeled antibody. The "two-step" assay involves washing before contacting the mixture with the labeled antibody. Other conventional methods may also be employed as suitable. It is usually desirable to immobilize one component of the assay system on a support, thereby allowing other components of the system to be brought into contact with the component and readily removed from the sample.

Streptococcus polypeptide-specific antibodies for use in the present invention can be raised against an intact *S. pneumoize* polypeptide of the present invention or fragment thereof. These polypeptides and fragments may be administered to an animal (*e.g.*, rabbit or mouse) either with a carrier protein (*e.g.*, albumin) or, if long enough (*e.g.*, at least about 25 amino acids), without a carrier.

As used herein, the term "antibody" (Ab) or "monoclonal antibody" (Mab) is meant to include intact molecules as well as antibody fragments (such as, for example, Fab and F(ab')₂ fragments) which are capable of specifically binding to a *Streptococcus* polypeptide. Fab and F(ab')₂ fragments lack the Fc fragment of intact antibody, clear more rapidly from the circulation, and may

have less non-specific tissue binding of an intact antibody (Wahl *et al.*, *J. Nucl. Med.* 24:316-325 (1983)). Thus, these fragments are preferred.

The antibodies of the present invention may be prepared by any of a variety of methods. For example, the *S. pneumoniae* polypeptide identified in Table 1, or fragments thereof, can be administered to an animal in order to induce the production of sera containing polyclonal antibodies. In a preferred method, a preparation of a *S. pneumoniae* polypeptide of the present invention is prepared and purified to render it substantially free of natural contaminants. Such a preparation is then introduced into an animal in order to produce polyclonal antisera of high specific activity.

In the most preferred method, the antibodies of the present invention are monoclonal antibodies. Such monoclonal antibodies can be prepared using hybridoma technology (Kohler *et al.*, *Nature* 256:495 (1975); Kohler *et al.*, *Eur. J. Immunol.* 6:511 (1976); Kohler *et al.*, *Eur. J. Immunol.* 6:292 (1976); Hammerling *et al.*, In: *Monoclonal Antibodies and T-Cell Hybridomas*, Elsevier, N.Y., (1981) pp. 563-681). In general, such procedures involve immunizing an animal (preferably a mouse) with a *S. pneumoniae* polypeptide antigen of the present invention. Suitable cells can be recognized by their capacity to bind anti-*Streptococcus* polypeptide antibody. Such cells may be cultured in any suitable tissue culture medium; however, it is preferable to culture cells in Earle's modified Eagle's medium supplemented with 10% fetal bovine serum (inactivated at about 56°C), and supplemented with about 10 g/l of nonessential amino acids, about 1,000 U/ml of penicillin, and about 100 µg/ml of streptomycin. The splenocytes of such mice are extracted and fused with a suitable myeloma cell line. Any suitable myeloma cell line may be employed in accordance with the present invention; however, it is preferable to employ the parent myeloma cell line (SP₂O), available from the American Type Culture Collection, Rockville, Maryland. After fusion, the resulting hybridoma cells are selectively maintained in HAT medium, and then cloned by limiting dilution as described by Wands *et al.* (*Gastroenterology* 80:225-232 (1981)). The hybridoma cells obtained through such a selection are then assayed to identify clones which secrete antibodies capable of binding the *Streptococcus* polypeptide antigen administered to immunized animal.

Alternatively, additional antibodies capable of binding to *Streptococcus* polypeptide antigens may be produced in a two-step procedure through the use of anti-idiotypic antibodies. Such a method makes use of the fact that antibodies are themselves antigens, and that, therefore, it is possible to obtain an antibody

which binds to a second antibody. In accordance with this method, *Streptococcus* polypeptide-specific antibodies are used to immunize an animal, preferably a mouse. The splenocytes of such an animal are then used to produce hybridoma cells, and the hybridoma cells are screened to identify clones which produce an antibody whose ability to bind to the *Streptococcus* polypeptide-specific antibody can be blocked by a *Streptococcus* polypeptide antigen. Such antibodies comprise anti-idiotypic antibodies to the *Streptococcus* polypeptide-specific antibody and can be used to immunize an animal to induce formation of further *Streptococcus* polypeptide-specific antibodies.

It will be appreciated that Fab and $F(ab')_2$ and other fragments of the antibodies of the present invention may be used according to the methods disclosed herein. Such fragments are typically produced by proteolytic cleavage, using enzymes such as papain (to produce Fab fragments) or pepsin (to produce $F(ab')_2$ fragments). Alternatively, *Streptococcus* polypeptide-binding fragments can be produced through the application of recombinant DNA technology or through synthetic chemistry.

Of special interest to the present invention are antibodies to *Streptococcus* polypeptide antigens which are produced in humans, or are "humanized" (*i.e.*, non-immunogenic in a human) by recombinant or other technology. Humanized antibodies may be produced, for example by replacing an immunogenic portion of an antibody with a corresponding, but non-immunogenic portion (*i.e.*, chimeric antibodies) (Robinson, R.R. *et al.*, International Patent Publication PCT/US86/02269; Akira, K. *et al.*, European Patent Application 184,187; Taniguchi, M., European Patent Application 171,496; Morrison, S.L. *et al.*, European Patent Application 173,494; Neuberger, M.S. *et al.*, PCT Application WO 86/01533; Cabilly, S. *et al.*, European Patent Application 125,023; Better, M. *et al.*, *Science* 240:1041-1043 (1988); Liu, A.Y. *et al.*, *Proc. Natl. Acad. Sci. USA* 84:3439-3443 (1987); Liu, A.Y. *et al.*, *J. Immunol.* 139:3521-3526 (1987); Sun, L.K. *et al.*, *Proc. Natl. Acad. Sci. USA* 84:214-218 (1987); Nishimura, Y. *et al.*, *Canc. Res.* 47:999-1005 (1987); Wood, C.R. *et al.*, *Nature* 314:446-449 (1985); Shaw *et al.*, *J. Natl. Cancer Inst.* 80:1553-1559 (1988). General reviews of "humanized" chimeric antibodies are provided by Morrison, S.L. (*Science*, 229:1202-1207 (1985)) and by Oi, V.T. *et al.*, *BioTechniques* 4:214 (1986)). Suitable "humanized" antibodies can be alternatively produced by CDR or CEA substitution (Jones, P.T. *et al.*, *Nature* 321:552-525 (1986);

Verhoeyan *et al.*, *Science* 239:1534 (1988); Beidler, C.B. *et al.*, *J. Immunol.* 141:4053-4060 (1988)).

Suitable enzyme labels include, for example, those from the oxidase group, which catalyze the production of hydrogen peroxide by reacting with substrate. Glucose oxidase is particularly preferred as it has good stability and its substrate (glucose) is readily available. Activity of an oxidase label may be assayed by measuring the concentration of hydrogen peroxide formed by the enzyme-labeled antibody/substrate reaction. Besides enzymes, other suitable labels include radioisotopes, such as iodine (^{125}I , ^{121}I), carbon (^{14}C), sulphur (^{35}S), tritium (^3H), indium (^{112}In), and technetium ($^{99\text{m}}\text{Tc}$), and fluorescent labels, such as fluorescein and rhodamine, and biotin.

Further suitable labels for the *Streptococcus* polypeptide-specific antibodies of the present invention are provided below. Examples of suitable enzyme labels include malate dehydrogenase, staphylococcal nuclease, delta-5-steroid isomerase, yeast-alcohol dehydrogenase, alpha-glycerol phosphate dehydrogenase, triose phosphate isomerase, peroxidase, alkaline phosphatase, asparaginase, glucose oxidase, beta-galactosidase, ribonuclease, urease, catalase, glucose-6-phosphate dehydrogenase, glucoamylase, and acetylcholine esterase.

Examples of suitable radioisotopic labels include ^3H , ^{111}In , ^{125}I , ^{131}I , ^{32}P , ^{35}S , ^{14}C , ^{51}Cr , ^{57}Co , ^{58}Co , ^{59}Fe , ^{75}Se , ^{152}Eu , ^{90}Y , ^{67}Cu , ^{217}Bi , ^{211}At , ^{212}Pb , ^{47}Sc , ^{109}Pd , etc. ^{111}In is a preferred isotope where *in vivo* imaging is used since it avoids the problem of dehalogenation of the ^{125}I or ^{131}I -labeled monoclonal antibody by the liver. In addition, this radionuclide has a more favorable gamma emission energy for imaging (Perkins *et al.*, *Eur. J. Nucl. Med.* 10:296-301 (1985); Carasquillo *et al.*, *J. Nucl. Med.* 28:281-287 (1987)). For example, ^{111}In coupled to monoclonal antibodies with 1-(P-isothiocyanatobenzyl)-DPTA has shown little uptake in non-tumorous tissues, particularly the liver, and therefore enhances specificity of tumor localization (Esteban *et al.*, *J. Nucl. Med.* 28:861-870 (1987)).

Examples of suitable non-radioactive isotopic labels include ^{157}Gd , ^{55}Mn , ^{162}Dy , ^{52}Tr , and ^{56}Fe .

Examples of suitable fluorescent labels include an ^{152}Eu label, a fluorescein label, an isothiocyanate label, a rhodamine label, a phycoerythrin label, a phycocyanin label, an allophycocyanin label, an o-phthaldehyde label, and a fluorescamine label.

Examples of suitable toxin labels include diphtheria toxin, ricin, and cholera toxin.

Examples of chemiluminescent labels include a luminal label, an isoluminal label, an aromatic acridinium ester label, an imidazole label, an acridinium salt label, an oxalate ester label, a luciferin label, a luciferase label, and an aequorin label.

5 Examples of nuclear magnetic resonance contrasting agents include heavy metal nuclei such as Gd, Mn, and iron.

Typical techniques for binding the above-described labels to antibodies are provided by Kennedy *et al.*, *Clin. Chim. Acta* 70:1-31 (1976), and Schurs *et al.*, *Clin. Chim. Acta* 81:1-40 (1977). Coupling techniques mentioned in the latter are the glutaraldehyde method, the periodate method, the dimaleimide method, the m-maleimidobenzyl-N-hydroxy-succinimide ester method, all of which methods are incorporated by reference herein.

10 In a related aspect, the invention includes a diagnostic kit for use in screening serum containing antibodies specific against *S. pneumoniae* infection. Such a kit may include an isolated *S. pneumoniae* antigen comprising an epitope which is specifically immunoreactive with at least one anti-*S. pneumoniae* antibody. Such a kit also includes means for detecting the binding of said antibody to the antigen. In specific embodiments, the kit may include a recombinantly produced or chemically synthesized peptide or polypeptide antigen. The peptide or polypeptide antigen may be attached to a solid support.

15 In a more specific embodiment, the detecting means of the above-described kit includes a solid support to which said peptide or polypeptide antigen is attached. Such a kit may also include a non-attached reporter-labelled anti-human antibody. In this embodiment, binding of the antibody to the *S. pneumoniae* antigen can be detected by binding of the reporter labelled antibody to the anti-*S. pneumoniae* antibody.

20 In a related aspect, the invention includes a method of detecting *S. pneumoniae* infection in a subject. This detection method includes reacting a body fluid, preferably serum, from the subject with an isolated *S. pneumoniae* antigen, and examining the antigen for the presence of bound antibody. In a specific embodiment, the method includes a polypeptide antigen attached to a solid support, and serum is reacted with the support. Subsequently, the support is reacted with a reporter-labelled anti-human antibody. The support is then examined for the presence of reporter-labelled antibody.

25 In a related aspect, the invention includes a method of detecting *S. pneumoniae* infection in a subject. This detection method includes reacting a body fluid, preferably serum, from the subject with an isolated *S. pneumoniae* antigen, and examining the antigen for the presence of bound antibody. In a specific embodiment, the method includes a polypeptide antigen attached to a solid support, and serum is reacted with the support. Subsequently, the support is reacted with a reporter-labelled anti-human antibody. The support is then examined for the presence of reporter-labelled antibody.

30 The solid surface reagent employed in the above assays and kits is prepared by known techniques for attaching protein material to solid support material, such as polymeric beads, dip sticks, 96-well plates or filter material. These attachment methods generally include non-specific adsorption of the

protein to the support or covalent attachment of the protein, typically through a free amine group, to a chemically reactive group on the solid support, such as an activated carboxyl, hydroxyl, or aldehyde group. Alternatively, streptavidin coated plates can be used in conjunction with biotinylated antigen(s).

Therapeutics and Modes of Administration

The present invention also provides vaccines comprising one or more polypeptides of the present invention. Heterogeneity in the composition of a vaccine may be provided by combining *S. pneumoniae* polypeptides of the present invention. Multi-component vaccines of this type are desirable because they are likely to be more effective in eliciting protective immune responses against multiple species and strains of the *Streptococcus* genus than single polypeptide vaccines. Thus, as discussed in detail below, a multi-component vaccine of the present invention may contain one or more, preferably 2 to about 20, more preferably 2 to about 15, and most preferably 3 to about 8, of the *S. pneumoniae* polypeptides identified in Table 1, or fragments thereof.

Multi-component vaccines are known in the art to elicit antibody production to numerous immunogenic components. Decker, M. and Edwards, K., *J. Infect. Dis.* 174:S270-275 (1996). In addition, a hepatitis B, diphtheria, tetanus, pertussis tetravalent vaccine has recently been demonstrated to elicit protective levels of antibodies in human infants against all four pathogenic agents. Aristegui, J. *et al.*, *Vaccine* 15:7-9 (1997).

The present invention thus also includes multi-component vaccines. These vaccines comprise more than one polypeptide, immunogen or antigen. An example of such a multi-component vaccine would be a vaccine comprising ~~more than one of~~ the *S. pneumoniae* polypeptides described in Table 1. A second example is a vaccine comprising ~~one or more, for example 2 to 10, of~~ the *S. pneumoniae* polypeptides identified in Table 1 and one or more, for example 2 to 10, additional polypeptides of either streptococcal or non-streptococcal origin. Thus, a multi-component vaccine which confers protective immunity to both a Streptococcal infection and infection by another pathogenic agent is also within the scope of the invention.

As indicated above, the vaccines of the present invention are expected to elicit a protective immune response against infections caused by species and strains of *Streptococcus* other than strain ~~of~~ *S. pneumoniae* deposited with that ATCC.

Further within the scope of the invention are whole cell and whole viral vaccines. Such vaccines may be produced recombinantly and involve the

expression of ~~one or more of~~ the *S. pneumoniae* polypeptides described in Table 1. For example, the *S. pneumoniae* polypeptides of the present invention may be either secreted or localized intracellular, on the cell surface, or in the periplasmic space. Further, when a recombinant virus is used, the *S. pneumoniae* polypeptides of the present invention may, for example, be localized in the viral envelope, on the surface of the capsid, or internally within the capsid. Whole cells vaccines which employ cells expressing heterologous proteins are known in the art. See, e.g., Robinson, K. *et al.*, *Nature Biotech.* 15:653-657 (1997); Sirard, J. *et al.*, *Infect. Immun.* 65:2029-2033 (1997); Chabalgoity, J. *et al.*, *Infect. Immun.* 65:2402-2412 (1997). These cells may be administered live or may be killed prior to administration. Chabalgoity, J. *et al.*, *supra*, for example, report the successful use in mice of a live attenuated *Salmonella* vaccine strain which expresses a portion of a platyhelminth fatty acid-binding protein as a fusion protein on its cells surface.

A multi-component vaccine can also be prepared using techniques known in the art by combining one or more *S. pneumoniae* polypeptides of the present invention, or fragments thereof, with additional non-streptococcal components (e.g., diphtheria toxin or tetanus toxin, and/or other compounds known to elicit an immune response). Such vaccines are useful for eliciting protective immune responses to both members of the *Streptococcus* genus and non-streptococcal pathogenic agents.

The vaccines of the present invention also include DNA vaccines. DNA vaccines are currently being developed for a number of infectious diseases. Boyer, J *et al.*, *Nat. Med.* 3:526-532 (1997); reviewed in Spier, R., *Vaccine* 14:1285-1288 (1996). Such DNA vaccines contain a nucleotide sequence encoding one or more *S. pneumoniae* polypeptides of the present invention oriented in a manner that allows for expression of the subject polypeptide. The direct administration of plasmid DNA encoding *B. burgdorgeri* OspA has been shown to elicit protective immunity in mice against borreliac challenge. Luke, C. *et al.*, *J. Infect. Dis.* 175:91-97 (1997).

The present invention also relates to the administration of a vaccine which is co-administered with a molecule capable of modulating immune responses. Kim, J. *et al.*, *Nature Biotech.* 15:641-646 (1997), for example, report the enhancement of immune responses produced by DNA immunizations when DNA sequences encoding molecules which stimulate the immune response are co-administered. In a similar fashion, the vaccines of the present invention may be co-administered with either nucleic acids encoding immune modulators or the immune modulators themselves. These immune modulators

include granulocyte macrophage colony stimulating factor (GM-CSF) and CD86.

5 The vaccines of the present invention may be used to confer resistance to streptococcal infection by either passive or active immunization. When the vaccines of the present invention are used to confer resistance to streptococcal infection through active immunization, a vaccine of the present invention is administered to an animal to elicit a protective immune response which either prevents or attenuates a streptococcal infection. When the vaccines of the present invention are used to confer resistance to streptococcal infection through
10 passive immunization, the vaccine is provided to a host animal (*e.g.*, human, dog, or mouse), and the antisera elicited by this antisera is recovered and directly provided to a recipient suspected of having an infection caused by a member of the *Streptococcus* genus.

15 The ability to label antibodies, or fragments of antibodies, with toxin molecules provides an additional method for treating streptococcal infections when passive immunization is conducted. In this embodiment, antibodies, or fragments of antibodies, capable of recognizing the *S. pneumoniae* polypeptides disclosed herein, or fragments thereof, as well as other *Streptococcus* proteins, are labeled with toxin molecules prior to their administration to the patient.
20 When such toxin derivatized antibodies bind to *Streptococcus* cells, toxin moieties will be localized to these cells and will cause their death.

The present invention thus concerns and provides a means for preventing or attenuating a streptococcal infection resulting from organisms which have antigens that are recognized and bound by antisera produced in
25 response to the polypeptides of the present invention. As used herein, a vaccine is said to prevent or attenuate a disease if its administration to an animal results either in the total or partial attenuation (*i.e.*, suppression) of a symptom or condition of the disease, or in the total or partial immunity of the animal to the disease.

30 The administration of the vaccine (or the antisera which it elicits) may be for either a "prophylactic" or "therapeutic" purpose. When provided prophylactically, the compound(s) are provided in advance of any symptoms of streptococcal infection. The prophylactic administration of the compound(s) serves to prevent or attenuate any subsequent infection. When provided
35 therapeutically, the compound(s) is provided upon or after the detection of symptoms which indicate that an animal may be infected with a member of the *Streptococcus* genus. The therapeutic administration of the compound(s) serves to attenuate any actual infection. Thus, the *S. pneumoniae* polypeptides, and

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fragments thereof, of the present invention may be provided either prior to the onset of infection (so as to prevent or attenuate an anticipated infection) or after the initiation of an actual infection.

5 The polypeptides of the invention, whether encoding a portion of a native protein or a functional derivative thereof, may be administered in pure form or may be coupled to a macromolecular carrier. Example of such carriers are proteins and carbohydrates. Suitable proteins which may act as macromolecular carrier for enhancing the immunogenicity of the polypeptides of the present invention include keyhole limpet hemacyanin (KLH) tetanus toxoid, 10 pertussis toxin, bovine serum albumin, and ovalbumin. Methods for coupling the polypeptides of the present invention to such macromolecular carriers are disclosed in Harlow *et al.*, *Antibodies: A Laboratory Manual, 2nd Ed.*; Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York (1988), the entire disclosure of which is incorporated by reference herein.

15 A composition is said to be "pharmacologically acceptable" if its administration can be tolerated by a recipient animal and is otherwise suitable for administration to that animal. Such an agent is said to be administered in a "therapeutically effective amount" if the amount administered is physiologically significant. An agent is physiologically significant if its presence results in a 20 detectable change in the physiology of a recipient patient.

While in all instances the vaccine of the present invention is administered as a pharmacologically acceptable compound, one skilled in the art would recognize that the composition of a pharmacologically acceptable compound varies with the animal to which it is administered. For example, a vaccine 25 intended for human use will generally not be co-administered with Freund's adjuvant. Further, the level of purity of the *S. pneumoniae* polypeptides of the present invention will normally be higher when administered to a human than when administered to a non-human animal.

30 As would be understood by one of ordinary skill in the art, when the vaccine of the present invention is provided to an animal, it may be in a composition which may contain salts, buffers, adjuvants, or other substances which are desirable for improving the efficacy of the composition. Adjuvants are substances that can be used to specifically augment a specific immune response. These substances generally perform two functions: (1) they protect 35 the antigen(s) from being rapidly catabolized after administration and (2) they nonspecifically stimulate immune responses.

Normally, the adjuvant and the composition are mixed prior to presentation to the immune system, or presented separately, but into the same

site of the animal being immunized. Adjuvants can be loosely divided into several groups based upon their composition. These groups include oil adjuvants (for example, Freund's complete and incomplete), mineral salts (for example, $\text{AlK}(\text{SO}_4)_2$, $\text{AlNa}(\text{SO}_4)_2$, $\text{AlNH}_4(\text{SO}_4)$, silica, kaolin, and carbon), polynucleotides (for example, poly IC and poly AU acids), and certain natural substances (for example, wax D from *Mycobacterium tuberculosis*, as well as substances found in *Corynebacterium parvum*, or *Bordetella pertussis*, and members of the genus *Brucella*. Other substances useful as adjuvants are the saponins such as, for example, Quil A. (Superfos A/S, Denmark). Preferred adjuvants for use in the present invention include aluminum salts, such as $\text{AlK}(\text{SO}_4)_2$, $\text{AlNa}(\text{SO}_4)_2$, and $\text{AlNH}_4(\text{SO}_4)$. Examples of materials suitable for use in vaccine compositions are provided in *Remington's Pharmaceutical Sciences* (Osol, A, Ed, Mack Publishing Co, Easton, PA, pp. 1324-1341 (1980), which reference is incorporated herein by reference).

The therapeutic compositions of the present invention can be administered parenterally by injection, rapid infusion, nasopharyngeal absorption (intranasopharangeally), dermoabsorption, or orally. The compositions may alternatively be administered intramuscularly, or intravenously. Compositions for parenteral administration include sterile aqueous or non-aqueous solutions, suspensions, and emulsions. Examples of non-aqueous solvents are propylene glycol, polyethylene glycol, vegetable oils such as olive oil, and injectable organic esters such as ethyl oleate. Carriers or occlusive dressings can be used to increase skin permeability and enhance antigen absorption. Liquid dosage forms for oral administration may generally comprise a liposome solution containing the liquid dosage form. Suitable forms for suspending liposomes include emulsions, suspensions, solutions, syrups, and elixirs containing inert diluents commonly used in the art, such as purified water. Besides the inert diluents, such compositions can also include adjuvants, wetting agents, emulsifying and suspending agents, or sweetening, flavoring, or perfuming agents.

Therapeutic compositions of the present invention can also be administered in encapsulated form. For example, intranasal immunization of mice against *Bordetella pertussis* infection using vaccines encapsulated in biodegradable microsphere composed of poly(DL-lactide-co-glycolide) has been shown to stimulate protective immune responses. Shahin, R. *et al.*, *Infect. Immun.* 63:1195-1200 (1995). Similarly, orally administered encapsulated *Salmonella typhimurium* antigens have also been shown to elicit protective

immunity in mice. Allaoui-Attarki, K. *et al.*, *Infect. Immun.* 65:853-857 (1997). Encapsulated vaccines of the present invention can be administered by a variety of routes including those involving contacting the vaccine with mucous membranes (*e.g.*, intranasally, intracolonicly, intraduodenally).

Many different techniques exist for the timing of the immunizations when a multiple administration regimen is utilized. It is possible to use the compositions of the invention more than once to increase the levels and diversities of expression of the immunoglobulin repertoire expressed by the immunized animal. Typically, if multiple immunizations are given, they will be given one to two months apart.

According to the present invention, an "effective amount" of a therapeutic composition is one which is sufficient to achieve a desired biological effect. Generally, the dosage needed to provide an effective amount of the composition will vary depending upon such factors as the animal's or human's age, condition, sex, and extent of disease, if any, and other variables which can be adjusted by one of ordinary skill in the art.

The antigenic preparations of the invention can be administered by either single or multiple dosages of an effective amount. Effective amounts of the compositions of the invention can vary from 0.01-1,000 $\mu\text{g/ml}$ per dose, more preferably 0.1-500 $\mu\text{g/ml}$ per dose, and most preferably 10-300 $\mu\text{g/ml}$ per dose.

Having now generally described the invention, the same will be more readily understood through reference to the following example which is provided by way of illustration, and is not intended to be limiting of the present invention, unless specified.

Examples

Example 1: Expression and Purification of *S. pneumoniae* Polypeptides in *E. coli*

The bacterial expression vector pQE10 (QIAGEN, Inc., 9259 Eton Avenue, Chatsworth, CA, 91311) is used in this example for cloning of the nucleotide sequence shown in Table 1 and for expressing the polypeptides identified in Table 1. The components of the pQE10 plasmid are arranged such that the inserted DNA sequence encoding a polypeptide of the present invention expresses the polypeptide with the six His residues (*i.e.*, a "6 X His tag") covalently linked to the amino terminus.

The DNA sequences encoding the desired portions of the polypeptides of Table 1 are amplified using PCR oligonucleotide primers from either a DNA

library constructed from *S. ⁴²pneumoniae*, such as the one deposited by the inventors at the ATCC for convenience, ATCC Deposit No. 97755, or from DNA isolated from the same organism such as the *S. pneumoniae* strain deposited with the ATCC as Deposit No. 55840. A list of PCR primers which can be used for this purpose is provided in Table 3, below. The PCR primers anneal to the nucleotide sequences encoding both the amino terminal and carboxy terminal amino acid sequences of the desired portion of the polypeptides of Table 1. Additional nucleotides containing restriction sites to facilitate cloning in the pQE10 vector were added to the 5' and 3' primer sequences, respectively. Such restriction sites are listed in Table 3 for each primer. In each case, the primer comprises, from the 5' end, 4 random nucleotides to prevent "breathing" during the annealing process, a restriction site (shown in Table 3), and approximately 15 nucleotides of *S. pneumoniae* ORF sequence (the complete sequence of each cloning primer is shown as SEQ ID NO: ³227 through SEQ ID NO: ⁴452).

For cloning the polypeptides of Table 1, the 5' and 3' primers were selected to amplify their respective nucleotide coding sequences. One of ordinary skill in the art would appreciate that the point in the protein coding sequence where the 5' primer begins may be varied to amplify a DNA segment encoding any desired portion of the complete amino acid sequences described in Table 1. Similarly, one of ordinary skill in the art would further appreciate that the point in the protein coding sequence where the 3' primer begins may also be varied to amplify a DNA segment encoding any desired portion of the complete amino acid sequences described in Table 1.

The amplified DNA fragment and the pQE10 vector are digested with the appropriate restriction enzyme(s) and the digested DNAs are then ligated together. The ligation mixture is transformed into competent *E. coli* cells using standard procedures such as those described in Sambrook *et al.*, *Molecular Cloning: a Laboratory Manual, 2nd Ed.*; Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y. (1989). Transformants are identified by their ability to grow under selective pressure on LB plates. Plasmid DNA is isolated from resistant colonies and the identity of the cloned DNA confirmed by restriction analysis, PCR and DNA sequencing.

Clones containing the desired constructs are grown overnight ("O/N") in liquid culture under selection. The O/N culture is used to inoculate a large culture, at a dilution of approximately 1:25 to 1:250. The cells are grown to an optical density at 600 nm ("OD600") of between 0.4 and 0.6. Isopropyl-b-D-thiogalactopyranoside ("IPTG") is then added to a final concentration of 1 mM

to induce transcription from the *lac* repressor sensitive promoter, by inactivating the *lacI* repressor. Cells subsequently are incubated further for 3 to 4 hours. Cells are then harvested by centrifugation. ^{4°C}

The cells are stirred for 3-4 hours at ~~4°C~~ in 6M guanidine-HCl, pH 8. The cell debris is removed by centrifugation, and the supernatant containing the protein of interest is loaded onto a nickel-nitrilo-tri-acetic acid ("NiNTA") affinity resin column (available from QIAGEN, Inc., *supra*). Proteins with a 6x His tag bind to the NI-NTA resin with high affinity and can be purified in a simple one-step procedure (for details see: The QIAexpressionist, 1995, QIAGEN, Inc., *supra*). Briefly, the supernatant is loaded onto the column in 6 M guanidine-HCl, pH8, the column is first washed with 10 volumes of 6 M guanidine-HCl, pH8, then washed with 10 volumes of 6 M guanidine-HCl pH6, and finally the polypeptide is eluted with 6 M guanidine-HCl, pH 5.0.

The purified protein is then renatured by dialyzing it against phosphate-buffered saline (PBS) or 50 mM Na-acetate, pH 6 buffer plus 200 mM NaCl. Alternatively, the protein can be successfully refolded while immobilized on the Ni-NTA column. The recommended conditions are as follows: renature using a linear 6M-1M urea gradient in 500 mM NaCl, 20% glycerol, 20 mM Tris/HCl pH7.4, containing protease inhibitors. The renaturation should be performed over a period of 1.5 hours or more. After renaturation the proteins can be eluted by the addition of 250 mM imidazole. Imidazole is removed by a final dialyzing step against PBS or 50 mM sodium acetate pH6 buffer plus 200 mM NaCl. The purified protein is stored at 4°C or frozen at -80°C.

The DNA sequences encoding the amino acid sequences of Table 1 may also be cloned and expressed as fusion proteins by a protocol similar to that described directly above, wherein the pET-32b(+) vector (Novagen, 601 Science Drive, Madison, WI 53711) is preferentially used in place of pQE10.

~~Each of the~~ ^{The} polynucleotides shown in Table 1, was successfully amplified and subcloned into pQE10 as described above using the PCR primers shown in Table 3. ~~These pQE10 plasmids containing the DNAs of Table 1,~~ except SP023, SP042, SP054, SP063, SP081, SP092, SP114, SP122, SP123, SP126, and SP127, were deposited with the ATCC as a pooled deposit as a convenience to those of skill in the art. This pooled deposit was deposited on October 16, 1997 and given ATCC Deposit No. 209369. Those of ordinary skill in the art appreciate that isolating an individual plasmid from the pooled deposit is trivial provided the information and reagents described herein. Each of the deposited clones is capable of expressing its encoded *S. pneumoniae* polypeptide.

Example 2: Immunization and Detection of Immune Responses

Methods

Growth of bacterial inoculum, immunization of Mice and Challenge with *S pneumoniae*.

Propagation and storage of, and challenge by *S. pneumoniae* are preformed essentially as described in Aaberge, I.S. et al., Virulence of *Streptococcus pneumoniae* in mice: a standardized method for preparation and frozen storage of the experimental bacterial inoculum, *Microbial Pathogenesis*, **18**:141 (1995), incorporated herein by reference.

Briefly, Todd Hewitt (TH) broth (Difco laboratories, Detroit, MI) with 17% FCS, and horse blood agar plates are used for culturing the bacteria. Both broth and blood plates are incubated at 37°C in a 5% CO₂ atmosphere. Blood plates are incubated for 18 hr. The culture broth is regularly 10-fold serially diluted in TH broth kept at room temperature and bacterial suspensions are kept at room temperature until challenge of mice.

For active immunizations C3H/HeJ mice (The Jackson Laboratory, Bar Harbor, ME) are injected intraperitoneally (i.p.) at week 0 with 20 g of recombinant streptococcal protein, or phosphate-buffered saline (PBS), emulsified with complete Freund's adjuvant (CFA), given a similar booster immunization in incomplete Freund's adjuvant (IFA) at week 4, and challenged at week 6. For challenge *S. pneumoniae* are diluted in TH broth from exponentially-growing cultures and mice are injected subcutaneously (s.c.) at the base of the tail with 0.1 ml of these dilutions (serial dilutions are used to find medium infectious dose). Streptococci used for challenge are passaged fewer than six times *in vitro*. To assess infection, blood samples are obtained from the distal part of the lateral femoral vein into heparinized capillary tubes. A 25 ul blood sample is serially 10-fold diluted in TH broth, and 25 ul of diluted and undiluted blood is plated onto blood agar plates. The plates are incubated for 18 hr. and colonies are counted.

Other methods are known in the art, for example, see Langermann, S. et al., *J. Exp. Med.*, **180**:2277 (1994), incorporated herein by reference.

Immunoassays

Several immunoassay formats are used to quantify levels of streptococcal-specific antibodies (ELISA and immunoblot), and to evaluate the functional properties of these antibodies (growth inhibition assay). The ELISA and immunoblot assays are also used to detect and quantify antibodies elicited in response to streptococcal infection that react with specific streptococcal antigens. Where antibodies to certain streptococcal antigens are elicited by infection this is taken as evidence that the streptococcal proteins in question are expressed *in vivo*. Absence of infection-derived antibodies (seroconversion) following streptococcal challenge is evidence that infection is prevented or suppressed. The immunoblot assay is also used to ascertain whether antibodies raised against recombinant streptococcal antigens recognize a protein of similar size in extracts of whole streptococci. Where the natural protein is of similar, or identical, size in the immunoblot assay to the recombinant version of the same protein, this is taken as evidence that the recombinant protein is the product of a full-length clone of the respective gene.

Enzyme-Linked Immunosorbant Assay (ELISA).

The ELISA is used to quantify levels of antibodies reactive with streptococcus antigens elicited in response to immunization with these streptococcal antigens. Wells of 96 well microtiter plates (Immunlon 4, Dynatech, Chantilly, Virginia, or equivalent) are coated with antigen by incubating 50 μ l of 1 μ g/ml protein antigen solution in a suitable buffer, typically 0.1 M sodium carbonate buffer at pH 9.6. After decanting unbound antigen, additional binding sites are blocked by incubating 100 μ l of 3% nonfat milk in wash buffer (PBS, 0.2% Tween 20, pH 7.4). After washing, duplicate serial two-fold dilutions of sera in PBS, Tween 20, 1% fetal bovine serum, are incubated for 1 hr, removed, wells are washed three times, and incubated with horseradish peroxidase-conjugated goat anti-mouse IgG. After three washes, bound antibodies are detected with H₂O₂ and 2,2'-azino-di-(3-ethylbenzthiazoline sulfonate) (Schwan, T.G., *et al.*, *Proc. Natl. Acad. Sci. USA* 92:2909-2913 (1985)) (ABTS®, Kirkegaard & Perry Labs., Gaithersburg, MD) and A₄₀₅ is quantified with a Molecular Devices, Corp. (Menlo Park, California) Vmax™ plate reader. IgG levels twice the background level in serum from naive mice are assigned the minimum titer of 1:100.

Sodiumdodecylsulfate-Polyacrylamide Gel Electrophoresis (SDS-PAGE) and Immunoblotting

Using a single well format, total streptococcal protein extracts or recombinant streptococcal antigen are boiled in SDS/2-ME sample buffer before electrophoresis through 3% acrylamide stacking gels, and resolving gels of higher acrylamide concentration, typically 10-15% acrylamide monomer. Gels are electro-blotted to nitrocellulose membranes and lanes are probed with dilutions of antibody to be tested for reactivity with specific streptococcal antigens, followed by the appropriate secondary antibody-enzyme (horseradish peroxidase) conjugate. When it is desirable to confirm that the protein had transferred following electro-blotting, membranes are stained with Ponceau S. Immunoblot signals from bound antibodies are detected on x-ray film as chemiluminescence using ECL™ reagents (Amersham Corp., Arlington Heights, Illinois).

Example 3: Detection of Streptococcus mRNA expression

Northern blot analysis is carried out using methods described by, among others, Sambrook *et al.*, *supra*. to detect the expression of the *S. pneumoniae* nucleotide sequences of the present invention in animal tissues. A cDNA probe containing an entire nucleotide sequence shown in Table 1 is labeled with ³²P using the *rediprime*™ DNA labeling system (Amersham Life Science), according to manufacturer's instructions. After labeling, the probe is purified using a CHROMA SPIN-100™ column (Clontech Laboratories, Inc.), according to manufacturer's protocol number PT1200-1. The purified labeled probe is then used to detect the expression of *Streptococcus* mRNA in an animal tissue sample.

Animal tissues, such as blood or spinal fluid, are examined with the labeled probe using ExpressHyb™ hybridization solution (Clontech) according to manufacturer's protocol number PT1190-1. Following hybridization and washing, the blots are mounted and exposed to film at -70 C overnight, and films developed according to standard procedures.

It will be clear that the invention may be practiced otherwise than as particularly described in the foregoing description and examples.

Numerous modifications and variations of the present invention are possible in light of the above teachings and, therefore, are within the scope of the appended claims.

The entire disclosure of all publications (including patents, patent applications, journal articles, laboratory manuals, books, or other documents) cited herein are hereby incorporated by reference.

E³

SP001 nucleotide (SEQ ID NO:1)

TAAAATCTACGACAATAAAAAATCAACTCATTTGCTGACTTGGGTTCTGAACGCCCGCTCAATGCCCAAGC
TAATGATATTCCACAGATTTGGTTAAGGCAATCGTTTCTATCGAAGACCATCGCTTCTTCGACCACAG
GGGGATTGATACCATCCGTATCCTGGGAGCTTTCTTGCAGCAATCTGCAAAGCAATTCCCTCCAAGGTGG
ATCAACTCTCACCAACAGTTGATTAAAGTTGACTTACTTTTCAACTTCGACTTCCGACCAGACTATTTC
TCGTAAGGCTCAGGAAGCTTGGTTAGCGATTGAGTTAGAACAAAAGCAACCAAGCAAGAAATCTTGAC
CTACTATATAAATAAGGTCTACATGTCTAATGGGAACATGGAATGCAGACAGCAGCTCAAACTACTA
TGGTAAAGACCTCAATAATTTAAGTTTACCTCAGTTAGCCTTGCTGGCTGGAATGCCTCAGCCACCAAA
CCAATATGACCCCTATTACATCCAGAAGCAGCCCAAGACCGCCGAAACTTGGTCTTATCTGAAATGAA
AAATCAAGGCTACATCTCTGCTGAACAGTATGAGAAAGCAGTCAATACACCAATTACTGATGGACTACA
AAGTCTCAAATCAGCAAGTAATTACCTTGCTTACATGGATAATTACCTCAAGGAAGTCAATCAATCAAGT
TGAAGAAGAAACAGGCTATAACCTACTCACAACCTGGGATGGATGTCTACACAAATGTAGACCAAGAAGC
TCAAAAACATCTGTGGGATATTTACAATACAGACGAATACGTTGCCTATCCAGACGATGAATTGCAAGT
CGCTTCTACCATTGTTGATGTTTCTAACGGTAAAGTCATTGCCAGCTAGGAGCAGCCCATCAGTCAAG
TAATGTTTCTTTCGGAATTAACCAAGCAGTAGAAACAAACCGCGACTGGGGATCAACTATGAAACCGAT
CACAGACTATGCTCCTGCCTTGGAGTACGGTGTCTACGATTCAACTGCTACTATCGTTCACGATGAGCC
CTATAACTACCCCTGGGACAAATACTCCTGTTTATAACTGGGATAGGGGCTACTTTGGCAACATCACCTT
GCAATACGCCCTGCAACAATCGCGAAACGTCACCGCGTGGAAACTCTAAACAAGGTCGGACTCGGACTCGG
CGCCAAGACTTTCTAAATGGTCTAGGAATCGACTACCCAAGTATTCACTACTCAAAATGCCATTTCAG
TAACACAACCGAATCAGACAAAAAATATGGAGCAAGTAGTGAAAAGATGGCTGCTGCTTACGCTGCCTT
TGCAATGGTGGAACTTACTATAAACCAATGTATATCCATAAAGTCTGCTTTAGTGATGGGAGTGAAAA
AGAGTTCTCTAATGTGCGAACTCGTGCCATGAAGGAAACGACAGCTATATGATGACCGACATGATGAA
AACAGTCTTGACTTATGGAACGGACGAAATGCCATCTTGCTTGGCTCCCTCAGGCTGGTAAAACAGG
AACCTCTAACTATACAGACGAGGAAATGAAAACCATCAAGACCTCTCAATTTGTAGCACCTGATGA
ACTATTTGCTGGCTATACGCGTAAATATCAATGGCTGTATGACAGGCTATTCTAACCGTCTGACACC
ACTTGTAGGCAATGGCCTTACGGTCGCTGCCAAAGTTTACCGCTCTATGATGACCTACCTGTCTGAAGG
AAGCAATCCAGAAGATTGGAATATACCAGAGGGGCTCTACAGAAATGGAGAATTCGATTTTAAAAATGG
TGCTCGTTCTACGTGGAACCTCACCTGCTCCACAACAACCCCATCAACTGAAAGTTCAAGTTCATCATC
AGATAGTTCAACTTCACAGTCTAGCTCAACCACTCCAAGCACAAATAATAGTACGACTACCAATCCTAA
CAATAATACGCAACAATCAAATACAACCCCTGATCAACAAAATCAGAATCCTCAACCAGCACAAACCA

SP001 AMINO ACID (SEQ ID NO:2)

KIYDNKNQLIADLGSERRVNAQANDIPTDLVKIVSIEDHRFFDHRGIDTIRILGAFLRNLSNSLQGG
STLTQQLIKLTYFSTSTSDQTI SRKAQEAWLHQLQEKATKQEILTYINKVYMSNGNYGMQTAAQNY
GKDLNLSLPQLALLAGMPQAPNQYDPYSHFEAAQDRRLVLSEMKNQGYISAEQYEKAVNTPITDGLQ
SLKSASNPAYMDNYLKEVINQVEEETGYMLLTGMDVYTNVDQEAQKHLWDIYNTDEVVAYPDDELQV
ASTIVDVSNGKVIAQLGARHQSSNVSFGLNQAVETNRDWGSTMKPITDYAPALEYGVYDSTATIVHDEP
YNYPGTNTPVYNWDRGYFGNITLQYALQSRNVPVAVETLNKVGLNRAKTFNLGLIDYPSIHYSNAISS
NTTESDKKYGASSEKMAAAYAAFANGSTYYKPMYIHKVVFSDGSEKEFSNVGTRAMKETAYMMTDMMK
TVLTYGTGRNAYLAWLPQAGKTGTSNYTDEEIHNIKTSQFVAPDELFAGYTRKYSMAVWTGYSNRLTP
LVGNGLTVAAKVYRSMMTYLSEGSNPEDWNIPEGLYRNGEFVKNGARSTWNSPAPQQPPSTESSSSSS
DSSTSQSSSTTPSTNNSTTTNPMNNTQQSNTPDQONQNPQPAQP

SP004 nucleotide (SEQ ID NO:3)

AAATTACAATACGGACTATGAATTGACCTCTGGAGAAAAATTACCTCTTCTCTAAAGAGATTTTCAGGTTA
CACTTATATTGGATATATCAAAGAGGGAAAAACGACTTCTGAGTCTGAAGTAAGTAATCAAAAAGAGTTC
AGTTGCCACTCCTACAAAACAACAAAAGGTGGATTATAATGTTACACCGAATTTTGTAGACCATCCATC
AACAGTACAAGCTATTTCAGGAACAAACACCTGTTTCTTCAACTAAGCCGACAGAAGTTCAAGTAGTTGA
AAAACCTTTCTCTACTGAATTAATCAATCCAAGAAAAGAAGAGAAACAATCTTCAGATTCTCAAGAACA
ATTAGCCGAACATAGAAATCTAGAAACGAAGAAAGAGGAGAAGATTTCTCCAAAAGAAAAGACTGGGGT
AAATACATTAAATCCACAGGATGAAGTTTATCAGGTCAATTGAACAAACCTGAACTCTTATATCGTGA
GGAAACTATGGAGACAAAAATAGATTTTCAAGAAGAAATCAAGAAAATCCTGATTTAGCTGAAGGAAC
TGTAAGAGTAAACAAGAAGGTAAATTAGTTAAGAAAGTTGAAATCGTCAGAATATTCTCTGTAAACAA
GGAAGAAGTTTCGCGAGAAATTGTTTCAACTTCAACGACTGCGCCTAGTCCAAGAATAGTCTGAAAAAGG
TACTAAAAAACTCAAGTTATAAAGGAACAACCTGAGACTGGTGTAGAACATAAGGACGTACAGTCTGG
AGCTATTGTTGAACCCGCAATTCAGCCTGAGTTGCCCGAAGCTGTAGTAAGTGACAAAGGCGAACCAGA
AGTTCAACCTACATTACCCGAAGCAGTTGTGACCGACAAAGGTGAGACTGAGGTTCAACCAGAGTCGCC
AGATACTGTGGTAAGTGATAAAGGTGAACCAGAGCAGGTAGCACCGCTTCCAGAATATAAGGGTAATAT

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Sub
E4

TGAGCAAGTAAAACCTGAAACTCCGGTTGAGAAGACCAAAGAACAAGGTCCAGAAAAAACTGAAGAAGT
TCCAGTAAAACCAACAGAAGAAACACCAGTAAATCCAAATGAAGGTACTACAGAAGGAACCTCAATTCA
AGAAGCAGAAAATCCAGTTCAACCTGCAGAAGAATCAACAACGAATTCAGAGAAAAGTATCACCAGATAC
ATCTAGCAAAAATACTGGGGAAGTGTCCAGTAATCCTAGTGATTGACAACCTCAGTTGGAGAATCAAA
TAAACCAGAACATAATGACTCTAAAAATGAAAATTCAGAAAAAACTGTAGAAGAAAGTTCCAGTAAATCC
AAATGAAGGCACAGTAGAAGGTACCTCAAATCAAGAAACAGAAAAACCAGTTCAACCTGCAGAAGAAAC
ACAAACAAACTCTGGGAAAATAGCTAACGAAAATACTGGAGAAGTATCCAATAAACCTAGTGATTCAAA
ACCACCAGTTGAAGAATCAAATCAACCAGAAAAAACGGAAGTCAACAAAAACCAGAAAAATTCAGGTAA
TACAACATCAGAGAATGGACAAACAGAACAGAACCATCAAACGGAATTCAACTGAGGATGTTTCAAC
CGAATCAAACACATCCAATTCAAATGGAACGAAGAAATTAACAAGAAAATGAAGTAGACCTGATAA
AAAGGTAGAAGAACAGAGAAAAACACTTGAATTAAGAAATGTTTCCGACCTAGAGTTA

SP004 amino acid (SEQ ID NO:4)

NYNTDYELTSGEKLPLPKEISGYTYIGYIKEGKTTSESEVSNQKSSVATPTKQQKVDYNVTPNFVDHPS
TVQAIQEQTTPVSSTKPTVEQVVEKPFSTELINPRKEEKQSSDSQEQLAEHKNLETKKEEKISPKEKTGV
NTLNPQDEVLSGQLNKPPELLYREETMETKIDFQEEIQENPDLAEGTVRVKQEGKLGKKVEIVRIFSVNK
EEVSREIVSTSTTAPSPRIVEKGTKKTKQVIKEQPETGVEHKDVQSGAIVEPAIQPELPEAVVSDKGEPE
VQPTLPEAVVTDKGETEVQPEPDTVSDKGEPEQVAPLPEYKGNIEQVKPETPVEKTKEQGPEKTEEV
PVKPTTEETPVNPNNEGTTGEGTSIQEAENPVQPAEESTTNSKVPSPDTSSKNTGEVSSNPSTSVGESN
KPEHNDSKNENSEKTVEEVPVNPNEGTVEGTSNQETEKPVQPAEETQTNSGKIANENTGEVSNKPSDSK
PPVEESNQPEKNGTATKPENSGNTTSENGQTEPEPSNGNSTEDVSTESNTSNSNGNEEIKQENELDPDK
KVEEPEKTLELRNVSDLEL

SP006 nucleotide (SEQ ID NO:5)

TGAGAATCAAGCTACACCCAAAGAGACTAGCGCTCAAAGACAATCGTCCTTGCTACAGCTGGCGACGT
GCCACCATTGACTACGAAGACAAGGGCAATCTGACAGGCTTTGATATCGAAGTTTAAAGGCAGTAGA
TGAAAACTCAGCGACTACGAGATTCAATTCCAAAGAACCGCCTGGGAGAGCATCTTCCCAGGACTTGA
TTCTGGTCACTATCAGGCTGCGGCCAATAACTTGAGTTACACAAAAGAGCGTGCTGAAAAATACCTTTA
CTCGCTTCCAATTTCCAACAATCCCCCTCGTCCTTGTCAGCAACAAGAAAAATCCTTTGACTTCTCTTGA
CCAGATCGCTGGTAAAACAACACAAGAGGATACCGGAACCTTCTAACGCTCAATTCATCAATAACTGGAA
TCAGAAACACACTGATAATCCCCTACAATTAATTTTTCTGGTGAGGATATTGGTAAACGAATCCTAGA
CCTTGCTAACGGAGAGTTTGATTTCCTAGTTTTTGTACAAGGTATCCGTTCAAAGATTATCAAGGACCG
TGGTTTAGACCTCTCAGTCGTTGATTACCTTCTGCAGATAGCCCCAGCAATTATATCATTTTCTCAAG
CGACCAAAAAGAGTTTAAAGAGCAATTTGATAAAGCGCTCAAAGAACTCTATCAAGACGGAACCTTGA
AAAACCTCAGCAATACCTATCTAGGTGGTTCTTACCTCCCAGATCAATCTCAGTTACAA

SP006 amino acid (SEQ ID NO:6)

ENQATPKETSQKTIIVLATAGDVPPFDYEDKGNLTGFDIEVLKAVDEKLSDYIEIQFORTAWESIFPGLD
SGHYQAAANNLSYTKERAELYSLPISNNPLVLVSNKKNPLTSLDQIAGKTTQEDTGTSNAQFINNWN
QKHTDNPATINFSGEDIGKRILDLANGEFDLFLVDKVSQKIIKDRGLDLSVVDLPSADSPSNYIIFSS
DQKEFKEQFDKALKELYQDGTLEKLSNTYLGGSYLPDQSQLQ

SP007 nucleotide (SEQ ID NO:7)

TGGTAACCGCTCTTCTCGTAACGCAGCTTCATCTTCTGATGTGAAGACAAAAGCAGCAATCGTCACTGA
TACTGGTGGTGTGATGACAAATCATTCAACCAATCAGCTTGGGAAGGTTTGCAGGCTTGGGGTAAAGA
ACACAATCTTTCAAAGATAACGGTTTCACTTACTTCCAATCAACAAGTGAAGCTGACTACGCTAACAA
CTTGCAACAAGCGGCTGGAAGTTACAACCTAATCTTCGGTGTGGTTTTGCCCTTAATAATGCAGTTAA
AGATGCAGCAAAAAGAACACACTGACTTGAACCTATGTCTTGATTGATGATGTGATTAAAGACCAAAAGAA
TGTTGCGAGCGTAACTTTTCGCTGATAATGAGTCAGGTTACCTTGCAGGTGTGGCTGCAGCAAAAACAAC
TAAGACAAAACAAGTTGGTTTTGTAGGTGGTATCGAATCTGAAGTTATCTCTCGTTTTGAAGCAGGATT
CAAGGCTGGTGTGCGTCAGTAGACCCATCTATCAAAGTCCAAGTTGACTACGCTGGTTCATTTGGTGA
TGCGGCTAAAGGTAAGGTAACCAATTGCAGCCGCACAATACGCAAGCCGGTGCAGATATTGTTTACCAAGTAGC
TGGTGGTACAGGTGCAGGTGTCTTTGCAGAGGCAAAATCTCTCAACGAAAGCCGCTCGTGAAAAATGAAAA
AGTTTGGGTATCGGTGTTGATCGTGACCAAGAAGCAGAAGGTAAATACACTTCTAAAGATGGCAAAGA
ATCAAACCTTGTCTTGTATCTACTTTGAAACAAGTTGGTACAACCTGTAAGATATTTCTAACAAAGGC
AGAAAGAGGAGAATTCCCTGGCGGTCAAGTGATCGTTTACTCATTGAAGGATAAAGGGGTTGACTTGGC
AGTAACAAACCTTTCAGAAGAAGGTAAAAAAGCTGTGCAAGATGCAAAAGCTAAAATCCTTGATGGAAG
CGTAAAGTTCTTGAAAAA

SP007 amino acid (SEQ ID NO:8)

GNRSSRNAASSSDVKTAAIVTDTGGVDDKSFNQSAWEGLQAWGKEHNLSKDNNGFTYFQSTSEADYANN
LQQAAGSYNLI FGVGFALNNAVKDAAKEHTDLNLYVLIDDDVIKDQKNVASVTFADNESGYLAGVAAAKTT
KTKQVGVFGGIESEVISRFEAGFKAGVASVDPISKVQVDYAGSFGDAAKGKTI AAAQYAAGADIVYQVA
GGTGAGVFAEAKSLNESRPENЕКVWVIGVDRDQEAEGKYTSKDGKESNFVLVSTLKQVGT TVKDISNKA
ERGEFPGGQVIVYSLKDKGVDLAVTNLSEEGKKAVEDAKAKILDG SVKVPEK

SP008 nucleotide (SEQ ID NO:9)

TGTGGAAATTTGACAGGTAACAGCAAAAAAGCTGCTGATTACAGGTGACAAACCTGTTATCAAAATGTAC
CAAATCGGTGACAAACCAGACAACCTTGGATGAATTGTTAGCAAATGCCAACAAAATCATTGAAGAAAAA
GTTGGTGCCAAATTGGATATCCAATACCTTGGCTGGGGTGACTATGGTAAGAAAATGTCAGTTATCACA
TCATCTGGTGAAAACCTATGATATTGCCTTTGCGAGATAACTATATTGTAAATGCTCAAAAAGGTGCTTAC
GCTGACTTGACAGAATTGTACAAAAAAGAAGGTAAAGACCTTTACAAAGCACTTGACCCAGCTTACATC
AAGGGTAATACTGTAAATGGTAAGATTTACGCTGTTCCAGTTGCAGCCAACGTTGCATCATCTCAAAAC
TTTGCCTTCAACGGAACCTCTCCTTGCTAAATATGGTATCGATATTTACAGGTGTTACTTCTTACGAACT
CTTGAGCCAGTCTTGAAACAAATCAAAGAAAAAGCTCCAGACGTAGTACCATTTGCTATTGGTAAAGTT
TTCATCCCCTCTGATAATTTGACTACCCAGTAGCAAAACGGTCTTCCATTGCTTATCGACCTTGAAGGC
GATACTACTAAAGTTGTAAACCGTTACGAAGTGCCTCGTTTCAAAGAACACTTGAAGACTCTTCACAAA
TTCTATGAAGCTGGCTACATTTCCAAAAGACGTCGCAACAAGCGATACTTCCCTTGACCTTCAACAAGAT
ACTTGGTTTCGTTTCGTGAAGAAACAGTAGGACCAGCTGACTACGGTAACAGCTTGCTTTTACGTTGCTTGC
AACAAAGATATCCAAATCAAACCAATTACTAATTCATCAAGNAAAACCAAACAACACAAGTTGCTAAC
TTTGTCTATCTCAAACAACCTTAAGAACAAAGAAAAATCAATGGAAATCTTGAACCTCTTGAATACGAAC
CCAGAACTCTTGAACGGTCTTGTTCACGGTCCAGAAGGCAAGAACTGGGAAAAAATTGAAGGTAAAGAA
AACCGTGTTCGCGTTCTTGATGGCTACAAAGGAAACACTCACATGGGTGGATGGAACACTGGTAACAAC
TGGATCCTTTACATCAACGAAAACGTTACAGACCAACAAATCGAAAATTCTAAGAAAGAATTGGCAGAA
GCTAAAGAATCTCCAGCGCTTGGAATTTATCTTCAATACTGACAATGTGAAATCTGAAATCTCAGCTATT
GCTAACACAATGCAACAATTTGATACAGCTATCAACACTGGTACTGTAGACCCAGATAAAGCGATTCCA
GAATTGATGGAAAAATTGAAATCTGAAGGTGCCTACGAAAAAGTATTGAACGAAATGCAAAAACAATAC
GATGAATTCTTGAAAAACAAAAAA

SP008 amino acid (SEQ ID NO:10)

CGNLTGNSKKAADSGDKPVIKMYQIGDKPDNLDELLANANKIIEEKVGAKLDIQYLGWGDYGKKMSVIT
SSGENYDIAFADNYIVNAQKGAYADLT ELYKKEGKDLYKALDPAYIKGNTVNGKIYAVPVAANVASSQN
FAFNGLLLAKYGIDISGVTSYETLEPVLKQIKEKAPDVVPFAIGKVFIPSDNFDYPVANGLPFVIDLEG
DTTKVVNRYEVPRFKEHLKTLHKFYEAGYIPKDVATSDTSFDLQQDTWFWVREETVGPADYNSLLSRVA
NKDIQIKPITNFIKXNQTTQVANFVISNNSKNKEKSMEILNLLNTNPELLNGLVYGPEGKNWEKIEGKE
NRVRVLDGYKGNTHMGWNTGNWILYINENVTDQQIENSKKELAEAKESPALGFIFNTDNVKSEISAI
ANTMQQFDTAINTGTVPDKAIPELMEKCLKSEGAYEKVLNEMQKQYDEFLKNKK

SP009 nucleotide (SEQ ID NO:11)

TGGTCAAGGAAGTCTTCTAAAGACAACAAAGAGGCAGAACTTAAGAAGGTTGACTTTATCCTAGACTG
GACACCAAATACCAACCACACAGGGCTTTATGTTGCCAAGGAAAAAGGTTATTTCAAAGAAGCTGGAGT
GGATGTTGATTTGAAATTGCCACCAGAAGAAAGTTCTTCTGACTTGATTATCAACGGAAAGGCACCATT
TGCAGTGTATTTCCAAGACTACATGGCTAAGAAATTGGAAAAAGGAGCAGGAATCACTGCCGTTGCAGC
TATTGTTGAACACAATACATCAGGAATCATCTCTCGTAAATCTGATAATGTAAGCAGTCCAAAAGACTT
GGTTGGTAAGAAATATGGGACATGGAATGACCCAACCTGAACCTTGCTATGTTGAAAACCTTGGTAGAATC
TCAAGGTGGAGACTTTGAGAAGGTTGAAAAAGTACCAAATAACGACTCAAACCTCAATCACACCGATTGC
CAATGGCGTCTTTGATACTGCTTGGAATTTACTACGGTTGGGATGGTATCCTTGCTAAATCTCAAGGTGT
AGATGCTAACTTCATGTACTTGAAAGACTATGTCAAGGAGTTTGACTACTATTACCAGTTATCATCGC
AAACAACGACTATCTGAAAGATAACAAAGAAGAAGCTCGCAAAGTCATCCAAGCCATCAAAAAAGGCTA
CCAATATGCCATGGAACATCCAGAAGAAGCTGCAGATATTCTCATCAAGAATGCACCTGAACCTCAAGGA
AAAACGTGACTTTGTCTCATCGAATCTCAAAAAATCTTGTGCAAAAGAATACGCAAGCGACAAGGAAAAATG
GGGTCAATTTGACGCAGCTCGCTGGAATGCTTTCTACAAATGGGATAAAGAAAAATGGTATCCTTAAAGA
AGACTTGACAGACAAAGGCTTCACCAACGAATTTGTGAAA

SP009 amino acid (SEQ ID NO:12)

GQGTASKDNKEAELKKVDFILDWTPNTNHTGLYVAKEKGYFKEAGVDVDLKLPPPESSSDLVINGKAPF
AVYFQDYMAKKLEKGAGITAVAAIVEHNTSGIISRKSDNVSSPKDLVGKKYGTWNDPTELAMLKTLVES
QGGDFEKVEKVPNNDSNSITPIANGVFDTAWIYYGWDGILAKSQGV DANFMYLKDYVKEFDYYSPIVIA
NNDYLKDNKEEARKVIOAIKKGYQYAMEHPPEEAADILIKNAPELKEKRDFVIESQKYLSKEYASDKEKW
GQFDAARWNAFYKWDKENGILKEDLTDKGFTNEFVK

SP010 nucleotide (SEQ ID NO:13)

TAGCTCAGGTGGAAACGCTGGTTCATCCTCTGGAAAAACAACCTGCCAAAGCTCGCACTATCGATGAAAT
CAAAAAAGCGGTGAACTGCGAATCGCCGTGTTTGGAGATAAAAAACCGTTTGGCTACGTTGACAATGA
TGGTTCTACCAAGGTACGCTACGATATTGAACTAGGGAACCAACTAGCTCAAGACCTTGGTGTCAAGGT
TAAATACATTTTCAGTCGATGCTGCCAACCGTGCGGAATACTTGATTTCAAACAAGGTAGATATTACTCT
TGCTAACTTTACAGTAACTGACGAACGTAAGAAACAAGTTGATTTTGCCCTTCCATATATGAAAGTTTC
TCTGGGTGTCGTATCACCTAAGACTGGTCTCATTACAGACGTCAAACAACCTTGAAGGTAAAAACCTTAAT
TGTCACAAAAGGAACGACTGCTGAGACTTATTTTGAAAAGAATCATCCAGAAATCAAACCTCCAAAAATA
CGACCAATACAGTGACTCTTACCAAGCTCTTCTTGACGGACGTGGAGATGCCTTTTCAACTGACAATAC
GGAAGTTCTAGCTTGGGCGCTTGAAAATAAAGGATTGAAGTAGGAATTACTTCCCTCGGTGATCCCGA
TACCATTGCGGCAGCAGTTCAAAAAGGCAACCAAGAATTGCTAGACTTCATCAATAAAGATATTGAAAA
ATTAGGCAAGGAAAACCTTCTTCCACAAGGCCTATGAAAAGACACTTCACCCAACCTACGGTGACGCTGC
TAAAGCAGATGACCTGGTTGTTGAAGGTGGAAAAGTTGAT

SP010 amino acid (SEQ ID NO:14)

SSGGNAGSSSGKTTAKARTIDEIKKSGELRIAVFGDKKPFYVDNDGSTKVRDYIELGNQLAQDLGVKV
KYISVDAANRAEYLISNKVDITLANFTVTDERKKQVDFALPYMKVSLGVVSPKTGLITDVKQLEGKTLI
VTKGTTAETYFEKNHPEIKLKQYDQYSDSYQALLDGRGDAFSTDNTEVLAWALENKGFEVGITSLGDPD
TIAAAVQKGNQELLDIFINKDIEKLGKENFFHKAYEKT LHPTYGDAAKADDLVVEGGKVD

SP011 nucleotide (SEQ ID NO:15)

CTCCAACATATGGTAAATCTGCGGATGGCACAGTGACCATCGAGTATTTCAACCAGAAAAAGAAATGAC
CAAAACCTTGGAAGAAATCACTCGTGATTTTGAGAAGGAAAACCCTAAGATCAAGGTCAAAGTCTGCAA
TGTAACCAATGCTGGTGAAGTATTGAAGACACGCGTTCGCGAGGAGATGTGCCTGATGTGGTCAATAT
TTACCCACAGTCCATCGAACTGCAAGAAATGGGCAAAAGCAGGTGTTTTTGAAGATTTGAGCAACAAAGA
CTACCTGAAACGCGTGAAAAATGGCTACGCTGAAAAATATGCTGTAAACGAAAAAGTTTACAACGTTCC
TTTTACAGCTAATGCTTATGGAATTTACTACAACAAAGATAAATTCGAAGAACTGGGCTTGAAGGTTCC
TGAAACCTGGGATGAATTTGAACAGTTAGTCAAAGATATCGTTGCTAAAGGACAAACACCATTTGGAAT
TGACAGGTGCAGATGCTTGGACACTCAATGGTTACAATCAATTAGCCTTTGCGACAGCAACAGGTGGAGG
AAAAGAAGCAAATCAATACCTTCGTTATTTCTCAACCAAATGCCATTAAATTGTCGGATCCGATTATGAA
AGATGATATCAAGGTCATGGACATCCTTCGCATCAATGGATCTAAGCAAAAGAACTGGGAAGGTGCTGG
CTATACCGATGTTATCGGAGCCTTCGCACGTGGGGATGTCTCATGACACCAAATGGGTCTTGGGCGAT
CACAGCGATTAAATGAACAAAAACCGAACTTTAAGATTGGGACCTTCATGATTCCAGGAAAAAGAAAAAGG
ACAAAGCTTAACCGTTGGTGCAGGAGACTTGGCATGGTCTATCTCAGCCACCACCAAACATCCAAAAGA
AGCCAATGCCTTTGTGGAATATATGACCCGTCCAGAAGTCATGCAAAAATACTACGATGTGGACGGATC
TCCAACAGCGATCGAAGGGGTCAAACAAGCAGGAGAAGATTACCGCTTGCTGGTATGACCGAATATGC
CTTTACGGATCGTCACTTGGTCTGGTTGCAACAATACTGGACCAGTGAAGCAGACTTCCATACCTTGAC
CATGAACATATGTCTTGACCGGTGATAAACAAGGCATGGTCAATGATTTGAATGCCTTCTTTAACCCGAT
GAAAGCGGATGTGGAT

SP011 amino acid (SEQ ID NO:16)

SNYGKSADGTVTIEYFNQKKEMTKLEEITRDFEKENPKIKVKVNVNPNAGEVLKTRVLAGDVPDVVNI
YPQSIELQEWAKAGVFEDLSNKDYLRVKNGYAEKYAVNEKVYNVPTANAYGIYYNKDFEELGLKVP
ETWDEFEQLVKDIVAKQTPFGIAGADAWTLNGYNQLAFATATGGGKEANQYLRYSQPNAIKLSDPIMK
DDIKVMDILRINGSKQKNWEGAGYTDVIGAFARGDVLMTPNGSWAITAINEQKPNFKIGTFMIPGKEKG
QSLTVGAGDLAWSISATTKHPKEANAFVEYMTREVMQKYVDVDSPTAIEGVKQAGEDSPLAGMTEYA
FTDRHLVWLQQYWTSEADFHTLTMNYVLTGDKQGMVNDLNAFFNPMKADVD

SP012 nucleotide (SEQ ID NO:17)

TGGGAAAAATCTAGCGAACTAGTGAGATAATTGGTCAAAGTACCAGTCTAACAAGTCTATTACTAT
TGGATTTGATAGTACTTTTGTTCGAATGGGATTTGCTCAGAAAGATGGTTCTTATGCAGGATTTGATAT
TGATTTAGCTACAGCTGTTTTTGAAAAATACGGAATCACGGTAAATTGGCAACCGATTGATTGGGATTT

GAAAGAAGCTGAATTGACAAAAGGAACGATTGATCTGATTTGGAATGGCTATTCCGCTACAGACGAACG
CCGTGAAAAGGTGGCTTTTCAGTAACCTCATATATGAAGAATGAGCAGGTATTGGTTACGAAGAAATCATC
TGGTATCACGACTGCAAAGGATATGACTGGAAAGACATTAGGAGCTCAAGCTGGTTTCATCTGGTTATGC
GGACTTTGAAGCAAATCCAGAAAATTTTGAAGAATATTGTCGCTAATAAGGAAGCGAATCAATACCAAAC
CTTTAATGAAGCCTTGATTGATTTGAAAAACGATCGAATTGATGGTCTATTGATTGACCGTGTCTATGC
AACTATTATTTAGAAGCAGAAGGTGTTTTAAACGATTATAATGTCTTTACAGTTGGACTAGAAACAGA
AGCTTTTTCGGTTGGAGCCCGTAAGGAAGATACAACTTGGTTAAGAAGATAAATGAAGCTTTTTCTAG
TCTTTACAAGGACGGCAAGTTCCAAGAAATCAGCCAAAAATGGTTTGGAGAAGATGTAGCAACCAAAGA
AGTAAAAGAAGGACAG

SP012 nucleotide (SEQ ID NO:18)

GKNSSETSGDNWSKYQSNKSITIGFDSFVPMGFAQKDGSYAGFDIDLATAVFKEYGITVNWQPIDWDL
KEAELTKGTIDLIWNGYSATDERREKVFASNSYMKNEQVLVTKSSGITTAKDMTGKTLGAQAGSSGYA
DFEANPEILKNIVANKEANQYQTFNEALIDLKNDRIDGLLIDRVYANYYLEAEGVLNDYNVFTVGLTE
AFAVGARKEDTNLVKKINEAFSSLYKDGKFQEIISQKWFGEDEVATKEVKEGQ

SP013 nucleotide (SEQ ID NO:19)

TGCTAGCGGAAAAAAGATACAACTTCTGGTCAAAAACATAAAGTTGTTGCTACAAACTCAATCATCGC
TGATATTACTAAAAATATTGCTGGTGACAAAATGACCTTCATAGTATCGTTCCGATTGGGCAAGACCC
ACACGAATACGAACCACTTCCTGAAGACGTTAAGAAAACCTCTGAGGCTAATTTGATTTCTATAACGG
TATCAACCTTGAAACAGGTGGCAATGCTTGGTTTACAAAATGGTAGAAAATGCCAAGAAAACCTGAAAA
CAAAGACTACTTCGCAGTCAGCGACGGCGTTGATGTTATCTACCTTGAAGGTCAAAAATGAAAAAGGAAA
AGAAGACCCACACGCTTGGCTTAACCTTGAAAACGGTATTATTTTTGCTAAAAATATCGCCAAACAATT
GAGCGCCAAAGACCCTAACAATAAAGAAATCTATGAAAAAATCTCAAAGAATATACTGATAAGTTAGA
CAAACCTTGATAAAGAAAGTAAGGATAAATTTAATAAGATCCCTGCTGAAAAGAAACTCATTGTAACCAG
CGAAGGAGCATTCAAATACTTCTCTAAAGCCTATGGTGTCCCAAGTGCTTACATCTGGGAAATCAATAC
TGAAGAAGAAGGAACCTCTGAACAAATCAAGACCTTGGTTGAAAACCTTCGCCAAACAAAAGTTCCATC
ACTCTTTGTAGAATCAAGTGTGGATGACCGTCCAATGAAAACCTGTTTCTCAAGACACAAACATCCCAAT
CTACGCTCAAATCTTTACTGACTCTATCGCAGAACAAAGGTAAAGAAGGCGACAGCTACTACAGCATGAT
GAAATACAACCTTGACAAGATTGCTGAAGGATTGGCAAAA

SP013 amino acid (SEQ ID NO:20)

ASGKKDTSYGQKLKVVATNSIIADITKNIAGDKIDLHSIVPIGQDPHEYEPLPEDVKKTSEANLIFYNG
INLETGGNAWFTKLVENAKKTENKDYFAVSDGVDVIYLEGQNEKGKEDPHAWLNLENGIIFAKNIAKQL
SAKDPNNKEFYEKNLKEYTDKLDKLDKESDKFNKI PAEKKLIVTSEGAFKYFSKAYGVPSAYIWEINT
EEEGTPEQIKTLVEKLRQTKVPSLFVSSVDDRPMKTVSQDTNIPYIAQIFTDSIAEQGKGDSYYSMM
KYNLDKIAEGLAK

SP014 nucleotide (SEQ ID NO:21)

TGGCTCAAAAATACAGCTTCAAGTCCAGATTATAAGTTGGAAGGTGTAACATTCCCGCTTCAAGAAAA
GAAAACATTGAAGTTTATGACAGCCAGTTCACCGTTATCTCCTAAAGACCCAAATGAAAAGTTAATTTT
GCAACGTTTGGAGAAGGAACTGGCGTTCATATTGACTGGACCAACTACCAATCCGACTTTGCAGAAAA
ACGTAACCTTGGATATTTCTAGTGGTGATTTACCAGATGCTATCCACAACGACGGAGCTTCAGATGTGGA
CTTGATGAACCTGGGCTAAAAAAGGTGTTATTATCCAGTTGAAGATTTGATTGATAAATACATGCCAAA
TCTTAAGAAAATTTTGGATGAGAAAACAGAGTACAAGGCCTTGATGACAGCACCTGATGGGCACATTTA
CTCATTTCCATGGATTGAAGAGCTTGGAGATGGTAAAGAGTCTATTACAGTGTCAACGATATGGCTTG
GATTAACAAAGATTGGCTTAAGAACTTGGTCTTGAAATGCCAAAACTACTGATGATTTGATTAAAGT
CCTAGAAGCTTTCAAAAACGGGGATCCAAATGGAATGGAGAGGCTGATGAAATTCATTTTTCATTTAT
TAGTGGTAACGGAAACGAAGATTTTAAATTCCTATTTGCTGCATTTGGTATAGGGGATAACGATGATCA
TTTAGTAGTAGGAAATGATGGCAAAGTTGACTTCACAGCAGATAACGATAACTATAAAGAAGGTGTCAA
ATTTATCCGTCAATTGCAAGAAAAAGGCCTGATTTGATAAAGAAGCTTTTCAACATGATTTGGAATAGTTA
CATTTGCTAAAGGTCATGATCAGAAAATTTGGTGTTTACTTTTACATGGGATAAGAATAAGTTACTGGAAG
TAACGAAAGTTATGATGTTTTTACCAGTACTTGTGTTGACCAAGTGGTCAAAAACACGTAGCTCGTACAAA
CGGTATGGGATTTGCACGTGACAAGATGGTTATTACCAGTGTAACAAAAACCTAGAATTGACAGCTAA
ATGGATTGATGCACAATACGCTCCACTCCAATCTGTGCAAAATAACTGGGGAACCTTACGGAGATGACAA
ACAACAAAACATCTTTGAATTGGATCAAGCGTCAAATAGTCTAAAACACTTACCCTAAACGGAACCTGC
ACCAGCAGAACTTCGTCAAAAGACTGAAGTAGGAGGACCCTAGCTATCCTAGATTACATACTATGGTAA
AGTAACAACCATGCCTGATGATGCCAAATGGCGTTTGGATCTTATCAAAGAATATTATGTCTCTTACAT

GAGCAATGTCAATAACTATCCAAGAGTCTTTATGACACAGGAAGATTTGGACAAGATTGCCCATATCGA
AGCAGATATGAATGACTATATCTACCGTAAACGTGCTGAATGGATTGTAAATGGCAATATTGATACTGA
GTGGGATGATTACAAGAAAGAACTTGAAAAATACGGACTTTCTGATTACCTCGCTATTAAACAAAAATA
CTACGACCAATACCAAGCAAACAAAAAC

SP014 amino acid (SEQ ID NO:22)

GSKNTASSPDYKLEGVTFPLQEKKTLKFMTASSPLSPKDPNEKLILQRLEKETGVHIDWTNYQSDFAEK
RNLDISSGDLPDIAIHNDGASVDLNMWAKKGVIIIPVEDLIDKYPNLLKKILDEKPEYKALMTAPDGHY
SFPWIEELGDGKESIHSVNDMAWINKDWLKKLGLEMPKTTDDLIKVLEAFKNGDPNGNGEADI PF SFI
SGNGNEDFKFLFAAFGIGDNDHLLVVGNDGKVDFTADNDNYKEGVKFI RQLQEKGLIDKEAFEHDWNSY
IAKGHDQKFGVYFTWDKNNVTGSNESYDVL PVLAGPSGQKHVARTNGMGFARDKMVITSVNKNLELTAK
WIDAQYAPLQSVQNNWGTYGDDKQONI FELDQASNSLKHLPNGTAPAELRQKTEVGGPLAILDSYYGK
VTMPDDAKWRLDLIKEYYVPYMSNVNYPVFMFTQEDLDKIAHIEADMNDYIYRKRAEWIVNGNIDTE
WDDYKKELEKYGLSDYLAIKQYYDQYQANKN

SP015 nucleotide (SEQ ID NO:23)

TAGTACAAACTCAAGCACTAGTCAGACAGAGACCAGTAGCTCTGCTCCAACAGAGGTAACCATTTAAAG
TTCATGGACGAGGTCAAACTTTCCAAAGTTCCTGAAAAGATTGTGACCTTTGACCTCGGCGCTGCGGA
TACTATTGCGCGCTTTAGGATTTGAAAAAATATCGTCGGAATGCCTACAAAAACTGTTCCGACTTATCT
AAAAGACCTAGTGGGAACGTCAAAAAATGTTGGTTCATGAAAGAACCTGATTTAGAAGCTATCGCCGC
CCTTGAGCCTGATTTGATTATCGCTTCGCCACGTACACAAAAATTCGTAGACAAATTCAAAGAAATCGC
CCCAACCGTTCTCTTCCAAGCAAGCAAGGACGACTACTGGACTTCTACCAAGGCTAATATCGAATCCCTT
AGCAAGTGCCTTCGGCGAAACTGGTACACAGAAAGCCAAGGAAGAATTGACCAAGCTAGACAAGAGCAT
CCAAGAAGTCGCTACTAAAAATGAAAGCTCTGACAAAAAGCCCTTGCGATCCTCCTTAATGAAGGAAA
AATGGCAGCCTTTGGTGCCAAATCTCGTTTCTCTTTCTTGTACCAAACCTTGAAATTCAAACCAACTGA
TACAAAATTTGAAGACTCACGCCACGGACAAGAAGTCAGCTTTGAAAGTGTCAAAGAAATCAACCCTGA
CATCCTCTTTGTTCATCAACCGTACCCTTGCCATCGGTGGGGACAACCTCTAGCAACGACGGTGTCTTAGA
AAATGCCCTTATCGCTGAAACACCTGCTGCTAAAAATGGTAAGATTATCCAACTAACACCAGACCTCTG
GTATCTAAGCGGAGGCGGACTTGAATCAACAAAACCTCATGATTGAAGACATACAAAAAGCTTTGAAA

SP015 amino acid (SEQ ID NO:24)

STNSSTSQTETSSSAPTEVTIKSSLDEVKLSKVPEKIVTFDLGAADTIRALGF EK NIVGMPTKTVPTYL
KDLVGT VKNV GSMKEPDLEAIAALEPDLIIASPR TQKFVDKFKEIAPT VLFQASKDDYWTSTKANIESL
ASAFGETGTQKAKEELTKLDKSIQEVATKNESD K KALAILLNEGKMAAFGA KSRFSFLYQTLKFKPTD
TKFEDSRHGQEVSFESVQKEINPDILFVINR TLAIGGDNSSNDGVLENALIAETPAKNGKIIQLTPDLW
YLSGGGLESTKLMIEDIQKALK

SP016 nucleotide (SEQ ID NO:25)

TGGCAATTCTGGCGGAAGTAAAGATGCTGCCAAATCAGGTGGTGACGGTGCCAAAACAGAAATCACTTG
GTGGGCATTCCCAGTATTTACCCAAGAAAAAACTGGTGACGGTGTTGGAACCTATGAAAAATCAATCAT
CGAAGCGTTTGAAAAAGCAAACCCAGATATAAAAGTGA AATTGGAAACCATCGACTTCAAGTCAGGTCC
TGAAAAAATCACAAACAGCCATCGAAGCAGGAACAGCTCCAGACGTACTCTTTGATGCACCAGGACGTAT
CATCCAATACGGTAAAAACGGTAAATTGGCTGAGTTGAATGACCTCTTCACAGATGAATTTGT TAAAGA
TGTCACAATGAAAACATCGTACAAGCAAGTAAAGCTGGAGACAAGGCTTATATGTATCCGATTAGTTC
TGCCCCATTCTACATGGCAATGAACAAGAAAAATGTTAGAAGATGCTGGAGTAGCAAACCTTGTA AAAAGA
AGGTTGGACAACCTGATGATTTTGAAAAAGTATTGAAAGCACTTAAAGACAAGGGTTACACACCAGGTTC
ATTGTTTCAGTTCTGGTCAAGGGGGAGACCAAGGAACACGTGCCTTTATCTCTAACCTTTATAGCGGTTC
TGTAACAGATGAAAAAGTTAGCAAATATACAACCTGATGATCCTAAATTCGTCAAAGGTCTTGAAAAAGC
AAGTAGCTGGATTAAAGACAATTTGATCAATAATGGTTTACAATTTGACGGTGGGGCAGATATCCAAAA
CTTTGCCAACGGTCAAACATCTTACACAATCCTTTGGGCACCAGCTCAAAATGGTATCCAAGCTAAACT
TTTAGAAGCAAGTAGAAGTGAAGTGGTAGAAGTACCATTCCCATCAGACGAAGGTAAGCCAGCTCTTGA
GTACCTTGTAACCGGTTTTCAGTATTCAACAATAAAGACGACAAGAAAGTCGCTGCATCTAAGAAAT
CATCCAGTTTATCGCAGATGACAAGGAGTGGGGACCTAAAGACGTAGTTTCGTACAGGTGCTTTCCAGT
CCGTACTTCATTTGAAAACTTTATGAAGACAAACGCATGGAAACAATCAGCGGCTGGACTCAATACTA
CTCACCATACTACAACACTATTGATGGATTTGCTGAAATGAGAACACTTTGGTTCCCAATGTTGCAATC
TGTATCAAATGGTGACGAAAAACCAGCAGATGCTTTGAAAGCCTTCACTGAAAAAGCGAACGAAACAAT
CAAAAAAGCTATGAAACAA

SP016 amino acid (SEQ ID NO:26)

GNSGGSKDAAKSGGDGAKTEITWWAFVFTQEKTDGVDGTYEKSIIIEAFEKANPDIKVKLETIDFKSGP
EKITTAIEAGTAPDVLFDAPGRIIQYGKNGKLAELNDLFTDEFVKDVNNENIVQASKAGDKAYMPISS
APFYMAMNKKMLEDAGVANLVKEGWTDDFEKVLKALKDKGYTPGSLFSSGQGGDQGTRAFISNLYSGS
VTDEKVSKEYTTDDPKFVKGLEKATSWIKDNLINNGSQFDGGADIQNFANGQTSYITLWAPAQNGIQAKL
LEASKVEVVEVPFSPDEGKPALEYLVNGFAVFNNKDDKKVAASKKFIQFIADDKEWGPKDVRTGAFPV
RTSFGKLYEDKRMETISGWTQYYSPYNTIDGFAEMRTLWF PMLQSVSNGDEKPADALKAFTEKANETI
KKAMKQ

SP017 nucleotide (SEQ ID NO:27)

TTCACAAGAAAAACAAAAAATGAAGATGGAGAACTAAGACAGAACAGACAGCCAAAGCTGATGGAAC
AGTCGGTAGTAAGTCTCAAGGAGCTGCCCAGAAGAAAGCAGAAGTGGTCAATAAAGGTGATTACTACAG
CATTCAGGGAAATACGATGAAATCATCGTAGCCAACAAACACTATCCATTGTCTAAAGACTATAATCC
AGGGGAAAATCCAACAGCCAAGGCAGAGTTGGTCAAACATCAAAAGCGATGCAAGAGGCAGGTTTCCC
TATTAGTGATCATTACAGTGGTTTTAGAAGTTATGAACTCAGACCAAGCTCTATCAAGATTATGTCAA
CCAGATGGAAAGGCAGCAGCTGACCGTTACTCTGCCCGTCTGGCTATAGCGAACACCAGACAGGCTT
GGCCTTTGATGTGATTGGGACTGATGGTGATTTGGTGACAGAAGAAAAGCAGCCCAATGGCTCTTGGA
TCATGCAGCTGATTATGGCTTTGTTGTCCGTTATCTCAAAGGCAAGGAAAAGGAAACAGGCTATATGGC
TGAAGAATGGCACCTGCGTTATGTAGGAAAAGAAGCTAAAGAAATTGCTGCAAGTGGTCTCAGTTTGGA
AGAATACTATGGCTTTGAAGCGGAGACTACGTCGAT

SP017 amino acid (SEQ ID NO:28)

SQEKTKNEDGETKTEQTAKADGTVGSKSQGAAQKKADEVVNKGDYYSIQGKYDEIIVANKHYPLSKDYNP
GENPTAKAELVKLIKAMQEAGFPISDHYSGRFSYETQTKLYQDYVNQDGKAAADRY SARPGYSEHQ TGL
AFDVIGTDGDLVTEEKAAQWLLDHAADYGFVVRYLKGKEKETGYMAEEWHLRYVGKEAKEIAASGLSLE
EYYGFEGGDYVD

SP019 nucleotide (SEQ ID NO:29)

GAAAGGTCTGTGGTCAAATAATCTTACCTGCGGTTATGATGAAAAATAATCTTGAAAAATATAAATAT
AAAAATACCTGAAGAAAAAATATCAGTTATTATTGGGTCAAATGGTTGTGGGAAATCAACACTCATTA
AACCTTGTCTCGACTTATAAAGCCATTAGAGGGAGAAGTATTGCTTGATAATAAATCAATTAATCTTA
TAAAGAAAAAGATTTAGCAAAACACATAGCTATATTACCTCAATCTCCAATAATCCCTGAATCAATAAC
AGTAGCTGATCTTGTAAAGCCGTGGTTCGTTTCCCTACAGAAAGCCTTTTAAGAGTCTTGAAAAAGATGA
CCTTGAAATAATAACAGATCAATGGTTAAGGCCAATGTTGAAGATCTAGCAAATAACCTAGTTGAAGA
ACTTTCTGGGGGTCAAAGGCAAAGAGTATGGATAGCTCTAGCCCTAGCCCAAGATACAAGTATCCTACT
TTTAGATGAGCCAACCTACTTGGATATCTCATATCAAATAGAACTATTAGACCTCTTGACTGATCT
AAACCAAAAATATAAGACAACCATTTGCATGATTTTGCACGATATAAATCTAACAGCAAGATACGCTGA
TTACCTATTTGCAATTAAAGAAGGTAAACTTGTGTCAGAGGGAAAGCCTGAAGATATACTAAATGATAA
ACTAGTTAAAGATATCTTTAATCTTGAAGCAAAAATTATACGTGACCCTATTTCCAATTCGCTCTAAT
GATTCCTATTGGCAAGCACCATGTTAACTCT

SP019 amino acid (SEQ ID NO:30)

KGLWSNNLTCGYDEKIILENINIKIPEEKISVIIGSNCGKSTLIKTL SRLIKPLEGEVLLDNKSINSY
KEKDLAKHIAILPQSPIIPESITVADLVSRGRFPYRKPFKSLGKDDLEIINRSMVKANVEDLANNLVEE
LSGGQRQRVWIALALAQDTSILLLLDEPTTYLDISYQIELDLLTDLNQKYKTTICMILHDINLTARYAD
YLF AIKEGKLVAEGKPEDILNDKLVKIDFNLEAKIIRDPISNSPLMIPIGKHHVS

SP020 nucleotide (SEQ ID NO:31)

AAACTCAGAAAAGAAAGCAGACAATGCAACAACCTATCAAAATCGCAACTGTTAACCGTAGCGGTTCTGA
AGAAAAACGTTGGGACAAAATCCAAGAATTGGTTAAAAAAGACGGAATTACCTTGGAATTTACAGAGTT
CACAGACTACTCACAAACCAACAAAGCAACTGCTGATGGCGAAGTAGATTGTAACGCTTTCCAACACTA
TAACCTTCTTGAACACTGGAACAAAGAAACGGAAGACCTTGTAGCGATTGCAGATACTTACATCTC
TCCAATCCGCCTTTACTCAGGTTTGAATGGAAGTGCCAACTAAAGTACACTAAAGTAGAAGACATCCCAGC
AAACGGAGAAATCGCTGTACCGAATGACGCTACAAACGAAAGCCGTGCGCTTTATTGCTTCAATCAGC
TGGCTTGATTAAATTGGATGTTTCTGGAAGTCTCTTGCAACAGTTGCCAACATCAAAGAAAAATCCAAA
GAACTTGAAAATCACTGAATTGGACGCTAGCCAAACAGCTCGTTTCAATTGTCATCAGTTGACGCTGCCGT
TGTAACAATACCTTCGTTACAGAAGCAAAATTGGACTACAAGAAATCACTTTTCAAAGAACAAGCTGA
TGAAAAC TCAAAACAATGGTACAACATCATTGTTGCAAAAAAGATTGGGAAACATCACCTAAGGCTGA

5.5

TGCTATCAAGAAAGTAATCGCAGCTTACCACACAGATGACGTGAAAAAAGTTATCGAAGAATCATCAGA
TGGTTTGGATCAACCAGTTTGG

SP020 amino acid (SEQ ID NO:32)

NSEKKADNATTIKIATVNRSGSEKRWKIQELVKKDGITLEFTEFTDYSQPNKATADGEVDLNAFQHY
NFLNNWNKENGKDLVAIADTYISPIRLYSGLNGSANKYTKVEDIPANGEIAVPNDATNESRALYLLQSA
GLIKLDVSGTALATVANIKENPKNLKITELDASQTARSLSSVDAAVVNNTFVTEAKLDYKKSFLKEQAD
ENSKQWYNIIVAKKDWETSPKADAIAIKVIAAYHTDDVKKVIEESSDGLDQPVW

SP021 nucleotide (SEQ ID NO:33)

TTCGAAAGGGTCAGAAGGTGCAGACCTTATCAGCATGAAAGGGGATGTCATTACAGAACATCAATTTTA
TGAGCAAGTGAAAAGCAACCCTTCAGCCCAACAAGTCTTGTTAAATATGACCATCCAAAAAGTTTGTGA
AAAACAATATGGCTCAGAGCTTGATGATAAAGAGGTTGATGATACTATTGCCGAAGAAAAAAAACAATA
TGGCGAAAACCTACCAACGTGTCTTGTCACAAGCAGGTATGACTCTTGAAACACGTAAAGCTCAAATTCG
TACAAGTAAATTAGTTGAGTTGGCAGTTAAGAAGGTAGCAGAAGCTGAATTGACAGATGAAGCCTATAA
GAAAGCCTTTGATGAGTACACTCCAGATGTAACGGCTCAAATCATCCGTCTTAATAATGAAGATAAGGC
CAAAGAAGTTCTCGAAAAAGCCAAGGCAGAAGGTGCTGATTTTGCTCAATTAGCCAAAGATAATTCAAC
TGATGAAAAAACAAAAGAAAATGGTGGAGAAATTACCTTTGATTCTGCTTCAACAGAAAGTACCTGGAGC
AAGTCCAAAAAAGCCGCTTTTCGCTTTTAGATGTGGGATGGTGTCTTCTGGATGTGGATTACAGCAACTG
GGGCACACCAAGCCTACAG

SP021 amino acid (SEQ ID NO:34)

SKGSEGADLISMKGDVITEHQFYEQVKSNPAAQVLLNMTIQKVFQYGSSELDDKEVDDTIAEEKQY
GENYQVRVLSQAGMTLETRKAQIRTSKLVELAVKKVAEAELEDEAYKKAFDEYTPDVTAQIIRLNEDKA
KEVLEKAKAEGADFAQLAKDNSTDEKTKENGGEITFDSASTEVPGASPKKPLFAFRCGMVFLDVDSNW
GTPSLQ

SP022 nucleotide (SEQ ID NO:35)

GGGGATGGCAGCTTTTAAAAATCCTAACAAATCAATACAAAGCTATTACAATTGCTCAAACTCTAGGTGA
TGATGCTTCTTCAGAGGAATTGGCTGGTAGATATGGTTCTGCTGTTTCAGTGACAGAAAGTGACTGCCTC
AAACCTTTCAACAGTTAAACTAAAGCTACGGTTGTAGAAAAACCACTGAAAGATTTTAGAGCGTCTAC
GTCTGATCAGTCTGGTTGGGTGGAATCTAATGGTAAATGGTATTTCTATGAGTCTGGTGATGTGAAGAC
AGGTTGGGTGAAAACAGATGGTAAATGGTACTATTTGAATGACTTAGGTGTCATGCAGACTGGATTTGT
AAAATTTTCTGGTAGCTGGTATTACTTGAGCAATTGAGGTGCTATGTTTACAGGCTGGGGAACAGATGG
TAGCAGATGGTTCTACTTTGACGGCTCAGGAGCTATGAAGACAGGCTGGTACAAGGAAAATGGCACTTG
GTATTACCTTGACGAAGCAGGTATCATGAAGACAGGTTGGTTTAAAGTCGGACCACACTGGTACTATGC
CTACGGTTCAGGAGCTTTGGCTGTGAGCACAAACACCAGATGGTTACCGTGTAATGGTAATGGTGA
ATGGGTAAAC

SP022 amino acid (SEQ ID NO:36)

GMAAFKNPNNQYKAITIAQTLGDDASSEELAGRYGSAVQCTEVTASNLSVKTATVVEKPLKDFRST
SDQSGWVESNGKWYFYESGDVKTGWVKTDGKWYYLNDLGVMQTGFVKFSGSWYYLSNSGAMFTGWGTDG
SRWFYFDGSGAMKTGWYKENGWYLLDEAGIMKTGWFKVGPWHYYAYGSGALAVSTTTPDGYRVNGNGE
WVN

SP023 nucleotide (SEQ ID NO:37)

AGACGAGCAAAAAATTAAGCAAGCAGAAGCGGAAGTTGAGAGTAAACAAGCTGAGGCTACAAGGTAA
AAAAATCAAGACAGATCGTGAAGAAGCAGAAGAAGAAGCTAAACGAAGAGCAGATGCTAAAGAGCAAGG
TAAACCAAAGGGGCGGGCAAAACGAGGAGTTCCCTGGAGAGCTAGCAACACCTGATAAAAAAGAAAATGA
TGCGAAGTCTTCAGATTCTAGCGTAGGTGAAGAACTCTTCCAAGCCCATCCCTGAAACCAGAAAAAA
GGTAGCAGAAGCTGAGAAGAAGGTTGAAGAAGCTAAGAAAAAGCCGAGGATCAAAAAGAAGAAGATCG
CCGTAACCTACCAACCAATACTTACAAAACGCTTGAACCTGAAATTGCTGAGTCCGATGTGGAAGTTAA
AAAAGCGGAGCTTGAAGTAGTAAAAAGAGGAAGCTAAGGAACCTCGAAACGAGGAAAAAGTTAAGCAAGC
AAAAGCGGAAGTTGAGAGTAAAAAAGCTGAGGCTACAAGGTTAGAAAAAATCAAGACAGATCGTAAAAA
AGCAGAAGAAGAAGCTAAACGAAAAAGCAGCAGAAGAAGATAAAGTTAAAGAAAAACCAGCTGAACAACC
ACAACCAGCGCCGGCTCCAAAAGCAGAAAAACCAGCTCCAGCTCCAAAACCAGAGAATCCAGCTGAACA
ACCAAAAGCAGAAAAACCAGCTGATCAACAAGCTGAAGAAGACTATGCTCGTAGATCAGAAGAAGAATA
TAATCGCTTGACTCAACAGCAACCGCCAAAAAATGAAAAACCAGCACAAACCATCTACTCCAAAAACAGG

CTGGAAACAAGAAAACGGTATGTGGTACTTCTACAATACTGATGGTTCAATGGCGACAGGATGGCTCCA
AAACAATGGCTCATGGTACTACCTCAACAGCAATGGCGCTATGGCGACAGGATGGCTCCAAAACAATGG
TTCATGGTACTATCTAAACGCTAATGGTTCAATGGCAACAGGATGGCTCCAAAACAATGGTTCATGGTA
CTACCTAAACGCTAATGGTTCAATGGCGACAGGATGGCTCCAATACAATGGCTCATGGTACTACCTAAA
CGCTAATGGTTCAATGGCGACAGGATGGCTCCAATACAATGGCTCATGGTACTACCTAAACGCTAATGG
TGATATGGCGACAGGTGGGTGAAAGATGGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAA
AGCAAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATGTCAATGGCTCAGGTGCCCTTGACGTCAA
CACAACGTAGATGGCTATGGAGTCAATGCCAATGGTGAATGGGTAAAC

SP023 amino acid (SEQ ID NO:38)

DEQKIKQAEAEVESKQAEATRLKKIKTDREEAEAEAKRRADAKEQGKPKGRAKRGVPGELATPDKKEND
AKSSDSSVGEETLPSPLKPEKKVAEAEKKVEEAKKAEDQKEEDRRNYPTNTYKTLLELEIAESDVEVK
KAELELVKEEAEKPRNEEKVKQAKAEVESKKAETRLKIKTDRKKAEEEEAKRKAEEEDKVKEKPAEQP
QPAPAPKAEPAPAPKPENPAEQPKAEKPADQQAEEEDYARRSEEEYNRLTQQQPPKTEKPAQPSTPKTG
WKQENGMWYFYNTDGSMTAGWLQNNGSWYYLNSNGAMATGWLQNNGSWYYLNANGSMATGWLQNNGSWY
YLNANGSMATGWLQYNGSWYYLNANGSMATGWLQYNGSWYYLNANGDMATGWVKDGDWYYLEASGAMK
ASQWFKVSDKWYYVNGSGALAVNTTVDGYGVNANGEWVN

SP025 nucleotide (SEQ ID NO:39)

CTGTGGTGAGGAAGAACTAAAAAGACTCAAGCAGCACAAACAGCCAAAACAACAAACGACTGTACAACA
AATTGCTGTTGGAAAAGATGCTCCAGACTTCACATTGCAATCCATGGATGGCAAAGAAGTTAAGTTATC
TGATTTTAAAGGGTAAAAAGGTTTACTTGAAGTTTGGGCTTCATGGTGTGGTCCATGCAAGAAAAGTAT
GCCAGAGTTGATGGAAGTAGCGGCGAAACCAGATCGTGATTTGGAATTCTTACTGTCAATTGCACCAGG
AATTCAAGGTGAAAAAAGTGTGAGCAATTCCACAATGGTTCCAGGAACAAGGATATAAGGATATCCC
AGTTCTTTATGATACCAAAGCAACCACTTCCAAGCTTATCAAATTCGAAGCATTCTTACAGAAATTT

SP025 amino acid (SEQ ID NO:40)

CGEEETKKTQAAQPKQQTTVQQIAVGKDAPDFTLQSMGKEVKLSDFKGGKVYLKFWASWCGPCKKSM
PELMELAAKPDRDFEILTVIAPGIQGEKTVQFPQWFQEQGYKDIPVLYDTKATTSKLIKFEAFQLNI

SP028 nucleotide (SEQ ID NO:41)

GACTTTTAAACAATAAACTATTGAAGAGTTGCACAATCTCCTTGTCTCTAAGGAAATTTCTGCAACAGA
ATTGACCCAAGCAACACTTGAAAAATATCAAGTCTCGTGAGGAAGCCCTCAATTCAATTTGTCACCATCGC
TGAGGAGCAAGCTCTTGTTCAAGCTAAAGCCATTGATGAAGCTGGAATTGATGCTGACAATGTCCTTTT
AGGAATTCCACTTGCTGTTAAGGATAACATCTCTACAGACGGTATTCTCACAACTGCTGCCTCAAAAAT
GCTCTACAACACTAGACCAATCTTTGATGCGCAGCTgTTGCCAATGCAAAAACCAAGGGCATGATTGT
CGTTGGAAAGACCAACATGGACGAATTTGCTATGGGTGGTTTCAGGtGAAACTTCACACTACGGAGCAAC
TAAAAACGCTTGGAACCAAGCAAGGTTCTGGTGGGTTCATCAAGTGGTTCTGCCGACGTGTAGCCTC
AGGACAAGTTCGCTTGTCACCTTGGTTCTGATACCTGGTGGTTCCATCCGCCAACCTGCTGCCTTCAACGG
AATCGTTGGTCTCAAAACCAACCTACGGAACAGTTTACGTTTTCGGTCTCATTGCCTTTGGTAGCTCATT
AGACCAGATTGGACCTTTTGCTCCTACTGTTAAGGAAATGCCCTCTTGCTCAACGCTATTGCCAGCGA
AGATGCTAAAGACTCTACTTCTGCTCCTGTCCGCATCGCCGACTTTACTTCAAAAATCGGCCAAGACAT
CAAGGGTATGAAAATCGCTTTGCCAAGGAATACCTAGGCGAAGGAATTGATCCAGAGGTTAAGGAAAC
AATCTTAAACGCGGCCAAACACTTTGAAAAATTTGGGTGCTATCGTCAAGAAGTCAGCCTTCCTCACTC
TAAATACGGTGTGGCGTTTATTACATCATCGCTTCATCAGAAGCTTCATCAAACCTGCAACGCTTCGA
CGGTATCCGTTACGGCTATCGCGCAGAAGATGCAACCAACCTTGATGAAATCTATGTAAACAGCCGAAG
CCAAGGTTTTGGTGAAGAGGTAAAACGTCGTATCATGCTGGGTACTTTTCAGTCTTTTCATCAGGTTACTA
TGATGCCTACTACAAAAGGCTGGTCAAGTCCGTACCTCATCATTCAAGATTTGCAAAAAGTCTTCGC
GGATTACGATTTGATTTTGGGTCCAAGTCTCAAGTGTTCCTATGACTTGGATTCTCTCAACCATGA
CCCAGTTGCCATGTACTTAGCCGACCTATTGACCATACCTGTAAACTTGGCAGGACTGCCTGGAATTTT
GATTCCTGCTGGATTCTCTCAAGGTCTACCTGTCCGACTCCAATTGATTGGTCCCAAGTACTCTGAGGA
AACCATTTACCAAGCTGCTGCTGCTTTTGAAGCAACAACAGACTACCACAAACAACAACCCGTGATTTT
TGGAGGTGACAAC

SP028 amino acid (SEQ ID NO:42)

TFNNKTIEELHNLLVSKEISATELTQATLENIKSREEALNSFVTIAEEQALVQAKAIDEAGIDADNVLS
GIPLAVKDNISTDGILTTAASKMLYNYEPIFDATAVANAKTKGMIVVGKTNMDEFAMGGSGETSHYGAT
KNAWNHKSVPGGSSSGSAAAVASGQVRLSLGSDTGGSIHQPAAFNGIVGLKPTYGTVSRFGLIAFGSSL

DQIGPFAPTVKENALLLNIAIASEDAKDSTSAPVRIADFTSKIGQDIKGMKIALPKKEYLGEIDPEVKET
ILNAAKHFEKLGAIVEEVSLPHSKYGVAVYIIASSEASSNLQRFDGIRYGYRAEDATNLDEIYVNSRS
QGFGEEVKRRIMLGTFSLSSGYDAYYKKAGQVRTLI IQDFEKVFADYDLILGPTAPSVAYDLDSLNDH
PVAMYLADLLTIPVNLAGLPGISIPAGFSQGLPVGLQLIGPKYSEETIYQAAAAFEATTDYHKQQPVIF
GGDN

SP030 nucleotide (SEQ ID NO:43)

CTTTACAGGTAAACAACACTACAAGTCGGCGACAAGGCGCTTGATTTTCTCTTACTACAACAGATCTTTC
TAAAAAATCTCTGGCTGATTTTGATGGCAAGAAAAAGTCTTGAGTGTCTTCCTTCTATCGATACAGG
CATCTGCTCAACTCAAACACGTCGTTTAAATGAAGAATTGGCTGGACTGGACAACACGGTCGTATTGAC
TGTTTCAATGGACCTACCTTTTGCTCAAAAACGTTGGTGCGGTGCTGAAGGCCTTGACAATGCCATTAT
GCTTTCAGACTACTTTGACCATTCTTTCGGGCGCGATTATGCCCTCTTGATCAACGAATGGCACCTATT
AGCACGCGCAGTCTTTGTCTCGATACTGACAATACGATTGCTACGTTGAATACGTGGATAATATCAA
TTCTGAGCCAAACTTCGAA

SP030 amino acid (SEQ ID NO:44)

FTGKQLQVGDKALDFSLTTTDL SKKSLADFDGKKKVL SVVPSIDTGICSTQTRRFNEELAGLDNTVVL
VSMDL PFAQKRWCGAEGLDNAIMLSDYFDHSFGRDYALLINEWHLLARAVFVLDTNTIRYVEYVDNIN
SEPNFE

SP031 nucleotide (SEQ ID NO:45)

CCAGGCTGATACAAGTATCGCAGACATTCAAAAAAGAGGCGAACTGGTTGTCGGTGTCAAAACAAGACGT
TCCCAATTTTGGTTACAAGATCCCAAGACCGGTACTTATTCTGGTATCGAAaCCGACTTGGCCAAGAT
GGTAGCTGATGAAGTCAAGGTCAAGATTCGCTATGTGCCGGTTACAGCACAAACCCGCGCCCCCTTCT
AGACAATGAACAGGTGATATGGATATCGCGACCTTTACCATCACGGACGAACGCAAAAACTCTACAA
CTTTACCAGTCCCTACTACACAGACGCTTCTGGATTTTGGTCAATAAATCTGCCAAAATCAAAAAGAT
TGAGGACCTAAACGGCAAAACCATCGGAGTCGCCCAAGGTTCTATCACCCAACGCCTGATTACTGAAGT
GGGTAAAAAGAAAGGTCTGAAGTTTAAATTCGTCGAACCTGGTTTCTACCCAGAATTGATTACTTCCCT
GCACGCTCATCGTATCGATACCTTTTCCGTTGACCGCTCTATTCTATCTGGCTACACTAGTAAACGGAC
AGCACTACTAGATGATAGTTTCAAGCCATCTGACTACGGTATTGTTACCAAGAAATCAAAATACAGAGCT
CAACGACTATCTTGATAACTTGGTTACTAAATGGAGCAAGGATGGTAGTTTGCAGAACTTTATGACCG
TTACAAGCTCAAACCATCTAGCCATACTGCAGAT

SP031 amino acid (SEQ ID NO:46)

QADTSIADIQKRGELVVGKQDVPNFGYXDPKGTGYSGIETDLAKMVADELKVKIRYVPVTAQTRGPLL
DNEQVMDIATFTITDERKKLYNFTSPYYTDASGFLVNKSAKIKKIEDLNGKTIGVAQGSITQRLITEL
GKKKGLKFKFVELGSYPELITSLHAHRIDTF SVDRSILSGYTSKRTALLDDSFKPSDYGIVTKKSNTL
NDYLDNLVTKWSKDGSLQKLYDRYKLKPSSTAD

SP032 nucleotide (SEQ ID NO:47)

GTCTGTATCATTTGAAAACAAAGAAACAAACCGTGGTGTCTTgACTTTCACTATCTCTCAAGACCAAAT
CAAACCAGAATTGGACCGTGTCTTCAAGtCAGTGAAGAAATCTCTTAATGTTCCAGGTTTCCGTAAAGG
TCACCTTCCACGCCCTATCTTCGACCAAAAATTTGGTGAAGAAGCTCTTTATCAAGATGCAATGAACGC
ACTTTTGCCAAACGCTTATGAAGCAGCTGTAAAAGAAGCTGGTCTTGAAGTGGTTGCCCAACCAAAAAT
TGACGTAACCTCAATGGAAAAAGGTCAAGACTGGGTATCACTGCTGAAGTCGTTACAAAACCTGAAGT
AAAATTTGGGTGACTACAAAACCTTGAAGTATCAGTTGATGTAGAAAAAGAAGTAACTGACGCTGATGT
CGAAGAGCGTATCGAACGCGAACGCAACAACCTGGCTGAATTGGTTATCAAGGAAGCTGCTGCTGAAAA
CGGCGACACTGTTGTGATCGACTTCGTTGGTCTATCGACGGTGTGAAATTTGACGGTGAAAAAGGTGA
AACTTCTCACTTGGACTTGGTTCAGGTCAATTCATCCCTGGTTTTCGAAGACCAATTGGTAGGTCACCTC
AGCTGGCGAAACCGTTGATGTTATCGTAACATTCCCAGAAGACTACCAAGCAGAAGACCTTGCAGGTAA
AGAAGCTAAATTCGTGACAACATCCACGAAGTAAAAGCTAAAGAAGTTCCGGCTCTTGACGATGAAT
TGCAAAAAGACATTGATGAAGAAGTTGAAACACTTGCTGACTTGAAAGAAAAATACAGCAAGAAGATTGGC
TGCTGCTAAAGAAGAAGCTTACAAAGATGCAGTTGAAGGTGCAGCAATTGATACAGCTGTAGAAAAATGC
TGAAATCGTAGAACTTCCAGAAGAAATGATCCATGAAGAAGTTCACCGTTCAGTAAATGAATTCCCTTGG
GAATTTGCAACGTCAAGGGATCAACCCTGACATGTACTTCCAAATCACTGGAACACTACTCAAGAAGACCT
TCACAACCAATACCAAGCAGAAGCTGAGTCACGTACTAAGACTAACCTTGTTATCGAAGCAGTTGCCAA
AGCTGAAGGATTTGATGCTTCAGAAGAAGAAATCCAAAAGAAGTTGAGCAATTGGCAGCAGACTACAA

CATGGAAGTTGCACAAGTTCAAAACTTGCTTTCAGCTGACATGTTGAAACATGATATCACTATCAAAAA
AGCTGTTGAATTGATCACAAGCACAGCAACAGTAAAA

SP032 amino acid (SEQ ID NO:48)

SVSFENKETNRGVLTFITISQDIKPELDRVFKSVKSLNVPGRKGLPRPIFDQKFGEEALYQDAMNA
LLPNAYEAAVKEAGLEVVAQPKIDVTSMEKGQDQWVITAEVVTKEVKLGDKYKNEVSVDVEKEVTDADV
EERIERERNLAEELVIKEAAAENGDTVIDFVGSIDGVEFDGGKGENFSLGLGSGQFIPGFEDQLVGH
AGETVDVIVTFPEDYQAEADLAGKEAKFVTTIHEVKAKEVPALDDELAKDIDEEVETLADLKEKYSKELA
AAKEEAYKDAVEGAIDTAVENAEIVELPEEMIHEEVHRSVNEFLGNLQROGINPDMYFQITGTTQEDL
HNQYQAEAESRTKTNLVIEAVAKAEGFDASEEEIQKEVEQLAADYNMEVAQVQNLASADMLKHDITIKK
AVELITSTATVK

SP033 nucleotide (SEQ ID NO:49)

TGGTCAAAAGGAAAGTCAGACAGGAAAGGGGATGAAAATTGTGACCAGTTTTTATCCTATCTACGCTAT
GGTTAAGGAAGTATCTGGTGACTTGAATGATGTTCCGATGATTCAGTCAAGTAGTGGTATTCACCTCTT
TGAACCTTCGGCAAATGATATCGCAGCCATCTATGATGCAGATGTCTTTGTTTACCATTCTCATACACT
CGAATCTTGGGCAGGAAGTCTGGATCCAAATCTAAAAAATCCAAAGTGAAGGTCTTAGAGGCTTCTGA
GGGAATGACCTTGGAAACGTGTCCCTGGACTAGAGGATGTGGAAGCAGGGGATGGAGTTGATGAAAAAC
GCTCTATGACCCTCACACATGGCTAGATCCTGAAAAGCTGGAGAAGAAGCCCAAATTATCGCTGATAA
ACTTTCAGAGGTGGATAGTGAGCATAAAGAGACTTATCAAAAAATGCGCAACCTTTATCAAAAAAGCT
CAGGAAT

SP033 amino acid (SEQ ID NO:50)

GQKESQTGKGMKIVTSFYPIYAMVKEVSGDLNDVRMIQSSSGIHSFEPSSANDIAAIYDADV FVYHSHTL
ESWAGSLDPNLKSKVKVLEASEGMTLERVPGLEDVEAGDGVDEKTLYPDHTWLDPEKAGEEAQIIADK
LSEVDSEHKETYQKNAQPLSKKLRLN

SP034 nucleotide (SEQ ID NO:51)

GAAGGATAGATATATTTTAGCATTTGAGACATCCTGTGATGAGACCAGTGTGCGCCGTCTTGAAAAACGA
CGATGAGCTCTTGTCCAATGTCAATGCTAGTCAAATTGAGAGTCACAAACGTTTTGGTGGCGTAGTGCC
CGAAGTAGCCAGTCGTCAACATGTGCGAGGTCAATACAGCCTGTATCGAGGAGGCATTGGCAGAAGCAGG
GATTACCGAAGAGGACGTGACAGCTGTTGCGGTTACCTACGGACCAGGCTTGGTGGAGCCTTGCTAGT
TGGTTTGTGCTGAGCTGCCAAGGCCTTTGCTTGGGCTCACGGACTTCCACTGATTCTCTGTTAATCACATGGC
TGGGCACCTCATGGCAGCTCAGAGTGTGGAGCCTTTGGAGTTTCCCTTGCTAGCCCTCTTGGTCAGCGG
CGGACACACAGAGTTGGTTTATGTTTTCGGAGCGGAGGATTATAAGATTGTTGGGGAAACCCGTGATGA
TGCGGTTGGTGAGGCTTATGATAAGGTGCGGCGTGTGATGGGCTTGACCTATCCTGCAGGTCGTGAGAT
TGACGAGCTGGCTCATCAGGGGCAGGATATTTATGATTTCCCCCGTGCCATGATTAAGGAAGATAATCT
GGAGTTCTCCTTCTCAGGTTTGAAATCTGCCTTTATCAATCTTCATCACAAATGCCGAGCAAAAAGGAGA
AAGCCTGTCTACAGAAGATTTGTGTGCTTCCCTTCCAAGCAGCAGTTATGGACATTCTCATGGCAAAAAC
CAAGAAGGCTTTGGAGAAATATCCTGTTAAATCCTAGTTGTGGCAGGTGGTGTGGCAGCCAATAAAGG
TCTCAGAGAACGCCTAGCAGCCGAAATCACAGATGTCAAGGTTATCATCCCCCTCTGCGACTCTGCGG
AGACAATGCAGGTATGATTGCCTATGCCAGCGTCAGCNAGTGAACAAAGAAAACCTTCGAGGCTGGGA
CCTCAATGCCAAACCAAGTCTTGCCCTTTGATACCATGGAA

SP034 amino acid (SEQ ID NO:52)

KDRYILAFETSCDETSVAVLKNDELLESNVIASQIESHKRFGGVVPEVASRHHVEVITACIEEALAEAG
ITEEDVTAVAVTYGPGLVGALLVGLSAAKAFWAHGLPLIPVNHMAGHLMAAQSVLEPFLALLVSG
GHTEL VYVSEAGDYKIVGETRDDAVGEAYDKVGRVMGLTYPAGREIDELAHQGDIDYDFPRAMIKEDNL
EFSFSGLKSAFINLHHNAEQKGESLSTEDLCASFQAAVMDILMAKTKKALEKYPVKILVVAGGVAAANKG
LRLRLAAEITDVKVIIPPLRLCGDNAGMIAYASVSXWNKENFAGWDLNAKPSLAFDTME

SP035 nucleotide (SEQ ID NO:53)

GGTAGTTAAAGTTGGTATTAACGTTTTCGGACGTATCGGTGCTCTTGCTTTCGGTCGTATCCAAAAACGT
AGAAGGTGTTGAAGTTACACGCATCAACGACCTTACAGATCCAGTTATGCTTGACACTTGTGAAAATA
CGACACAACCTCAAGGTCGTTTTCGACGGTACTGTTGAAGTTAAAGAAGGTGGATTTGAAGTTAACGGTAA
ATTCATCAAAGTTTCTGCTGAACGTGATCCAGAACAATCGACTGGGCTACTGACGGTGTAGAAATCGT
TCTTGAAGCTACTGGTTTCTTTGCTAAGAAAGAAGCAGCTGAAAAACACCTTAAAGGTGGAGCTAAAAA

AGTTGTTATCACTGCTCCTGGTGGAACGACGTTAAACAGTTGTATTCAACACTAACCACGACGTTCT
TGACGGTACTGAAACAGTTATCTCAGGTGCTTCATGTACTACAACTGCTTGGCTCCAATGGCTAAAGC
TCTTCAAGACAACCTTTGGTGTGTTGAAGGATTGATGACTACTATCCACGCTTACACTGGTGACCAAAT
GATCCTTGACGGACCACACCGTGGTGGTGACCTTCGCCGTGCTCGCGCTGGTGCTGCAAAACATCGTTCC
TAACCTCAACTGGTGCTGCAAAAGCTATCGGTCTTGTAATCCCAGAATTGAATGGTAAACTTGACGGATC
TGCACAACGCGTTCCAACCTCCAACCTGGATCAGTTACTGAATTGGTAGCAGTTCTTGAAAAGAAGCTTAC
TGTTGATGAAGTGAACGCAGCTATGAAAGCAGCTTCAAACGAATCATACGGTTACACAGAAGATCCAAT
CGTATCTTCAGATATCGTAGGTATGTCTTACGGTTCATTGTTTGACGCAACTCAAACCTAAAGTTCTTGA
CGTTGACGGTAAACAATTGGTTAAAGTTGTATCATGGTACGACAACGAAATGTCATACACTGCACAAC
TGTTCTGACTCTTGAATACTTCGCAAAAATTGC

SP035 amino acid (SEQ ID NO:54)

VVKVINGIFGRIGRLAFRRRIQNVEGVEVTRINDLTDPMVLAHLLKYDTTQGRFDGTVEVKEGGFEVNGK
FIKVSARDEPEQIDWATDGEIVLEATGFFAKKEAAEKHLKGGAKKVITAPGGNDVKTVVFNTNHDVL
DGTETVISGASCTTNCLAPMAKALQDNFGVVEGLMTTIIHAYTGDQMILDGPHRGGDLRRARAGANIVP
NSTGAAKAIGLVIPELNGKLDGSAQRVPTPTGSVTELAVLEKNVTVDEVNAAMKAASNESYGYTEDPI
VSSDIVGMSYGSFLDATQTKVLDVDGKQLVKVSWYDNEMSYTAQLVRTLGLRKNC

SP036 nucleotide (SEQ ID NO:55)

TTCTTACGAGTTGGGACTGTATCAAGCTAGAACGGTTAAGGAAAATAATCGTGTTTCCTATATAGATGG
AAAACAAGCGACGCAAAAAACGGAGAATTTGACTCCTGATGAGGTTAGCAAGCGTGAAAGGAATCAATGC
TGAGCAAATCGTCATCAAGATAACAGACCAAGGCTATGTCACCTCACATGGCGACCACTATCATTATTA
CAATGGTAAGGTTTCCTTATGACGCTATCATCAGTGAAGAATTACTCATGAAAGATCCAAACTATAAGCT
AAAAGATGAGGATATTGTTAATGAGGTCAAGGTTGATATGTTATCAAGGTAGATGGAAAATACTATGT
TTACCTTAAGGATGCTGCCCACGCGGATAACGTCCTGACAAAAGAGGAAATCAATCGACAAAAACAAGA
GCATAGTCAACATCGTGAAGGTGGAACCTCAAAGAACGATGGTGCTGTTGCCTTGGCACGTTTCGCAAGG
ACGCTATACACTACAGATGATGGTTATATCTTTAATGCTTCTGATATCATAGAGGATACTGGTGATGCTTA
TATCGTTTCCTCATGGAGATCATTACCATTACATTCCTAAGAATGAGTTATCAGCTAGCGAGTTGGCTGC
TGCAGAAGCCTTCCTATCTGGTTCGAGGAAAATCTGTCAAATTCAGAACCTATCGCCGACAAAATAGCGA
TAACACTTCAAGAACAACTGGGTACCTTCTGTAAGCAATCCAGGAACCTACAAATACTAACACAAGCAA
CAACAGCAACACTAACAGTCAAGCAAGTCAAAGTAATGACATTGATAGTCTCTTGAAACAGCTCTACAA
ACTGCCCTTTGAGTCAACGACATGTAGAATCTGATGGCCTTGTCTTTGATCCAGCACAAATCACAAGTCG
AACAGCTAGAGGTGTTGCAGTGCCACACGGAGATCATTACCACTTCATCCCTTACTCTCAAATGTCTGA
ATTGGAAGAACGAATCGCTCGTATTATCCCTTTCGTTTCAAAACCTTGGGTACCAGATTCAAAG
GCCAGAACAAACCAAGTCCACAACCGACTCCGGAACCTAGTCCAGGCCCGCAACCTGCACCAATCTTAA
AATAGACTCAAATTCCTTCTTTGGTTAGTCAGCTGGTACGAAAAGTTGGGGAAGGATATGTTATCGAAGA
AAAGGGCATCTCTCGTTATGTCTTTGCGAAAAGATTTACCATCTGAAACTGTTAAAAATCTTGAAAGCAA
GTTATCAAAACAAGAGAGTGTTCACACACTTTAACTGCTAAAAAAGAAAATGTTGCTCCTCGTGACCA
AGAATTTTATGATAAAGCATATAATCTGTAACTGAGGCTCATAAAGCCTTGTTTGNAAAATAAGGGTCG
TAATCTGATTTCCAAGCCTTAGACAAATTAATAGAACGCTTGAATGATGAATCGACTAATAAAGAAAA
ATTGGTAGATGATTTATTGGCATTCTTAGCACCAATTACCCATCCAGAGCGACTTGGCAAACCAAATTC
TCAAATTGAGTATACTGAAGACGAAGTTCGTATTGCTCAATTAGCTGATAAGTATACAACGTCAGATGG
TTACATTTTTTGATGAACATGATATAATCAGTGATGAAGGAGATGCATATGTAACGCCCTCATATGGGCCA
TAGTCACTGGATTGGAAAAGATAGCCTTCTGATAAGGAAAAGTTGCAGCTCAAGCCTATACATAAGA
AAAAGGTATCCTACCTCCATCTCCAGACGCAGATGTTAAAGCAAATCCAACCTGGAGATAGTGCAGCAGC
TATTTACAATCGTGTGAAAGGGGAAAAACGAATTCACCTCGTTCGACTTCCATATATGGTTGAGCATAC
AGTTGAGGTTAAAAACGGTAATTTGATTATTCCTCATAAGGATCATTACCATAATATTAAATTTGCTTG
GTTTGATGATCACACATACAAAGCTCCAAATGGCTATACCTTGAAGATTGTTTTCGACGATTAAAGTA
CTACGTAGAACACCTTGACGAACGTCCACATCTAATGATGGATGGGGCAATGCCAGTGAGCATGTGTT
AGGCAAGAAAGACCACAGTGAAGATCCAAATAAGAACTTCAAAGCGGATGAAGAGCCAGTAGAGGAAAC
ACCTGCTGAGCCAGAAGTCCCTCAAGTAGAGACTGAAAAGTAGAAGCCCAACTCAAAGAAGCAGAAGT
TTTGCTTGCGAAAAGTAACGGATTCTAGTCTGAAAGCCCAATGCAACAGAACTCTAGCTGGTTTACGAAA
TAATTTGACTCTTCAAATTATGGATAACAATAGTATCATGGCAGAAGCAGAAAAATTACTTGCGTTGTT
AAAAGGAAGTAATCCTTCATCTGTAAGTAAGGAAAAAATAAAC

SP036 amino acid (SEQ ID NO:56)

SYELGLYQARTVKENNRVSYIDGKQATQKTENLTPDEVSKREGINAEQIVIKITDQGYVTSBGDHYHY
NGKVPYDAIISELLMKDPNYKLKDEDIVNEVKGGYVIKVDGKYYVYLKDAHADNVRTKEEINRQKQE

260907-10999680

HSQHREGGTPRNDGAVALARSQGRYTTDDGYIFNASDIIEDTGDAYIVPHGDHYHYIPKNELSASELAA
AEAFLSGRGNLSNSRTYRRQNSDNTSRTNWVPSVSNPGTTNTNTSNNSTNSQASQSNDDIDSLKQLYK
LPLSQRHVESDGLVFDPAQITSRTARGVAVPHGDHYHFIPYSQMSELEERIARIIPLYRSNHWVPDSR
PEQSPQPTPEPSPGPQPAPNLKIDSSSLVSQLVRKVGEGYVFEEKGISRYVFAKDLPSSETVKNLESK
LSKQESVSHTLTAKKENVAPRDQEFYDKAYNLLTEAHKALFXNKGNSDFQALDKLLERLNDESTNKEK
LVDDLLAFLAPITHPERLGKPNSEQIEYTEDEVRIAQLADKYTTSDGYIFDEHDIISDEGDAYVTPHMGH
SHWIGKDSLSDEKVAQAAYTKEKGILPPSPDADVKNAPTGDSAAAIYNRVKGEKRIPLVRLPYMVEHT
VEVKNGNLIIPHKDHYHNIKFAWFDDHTYKAPNGYTLEDLFATIKYYVEHPDERPHSNDGWGNASEHVL
GKKDHSSEDPNKNFKADEEPVEETPAEPEVPQVETEKVEAQLKEAEVLLAKVTDSSSLKANATETLAGLRN
NLTLQIMDNNSIMAEAEKLLALLKGSNPSSVSKEKIN

SP038 nucleotide (SEQ ID NO:57)

TACTGAGATGCATCATAATCTAGGAGCTGAAAAGCGTTTCAGCAGTGGCTACTACTATCGATAGTTTTAA
GGAGCGAAGTCAAAAAGTCAGAGCACTATCTGATCCAAATGTGCGTTTTGTTCCCTTCTTTGGCTCTAG
TGAATGGCTTCGTTTTGACGGTGCTCATCTGCGGTATTAGCTGAGAAATACAATCGTTCCTACCGTCC
TTATCTTTTAGGACAGGGGGGAGCTGCATCGCTTAACCAATATTTTGAATGCAACAGATGTTACCACA
GCTGGAGAATAAAACAAGTTGTGTATGTTATCTCACCTCAGTGGTTCAGTAAAAATGGCTATGATCCAGC
AGCCTTCCAGCAGTATTTTAATGGAGACCACTGACTACTTTTCTGAAACATCAATCTGGGGATCAGGC
TAGTCAATATGCAGCGACTCGCTTACTGCAACAGTTCCCAAACGTAGCTATGAAGGACCTGGTTAGAA
GTTGGCAAGTAAAGAAGAATTGTCGACGACGACAATGAAATGATTGAATTATTGGCTCGTTTTTAATGA
ACGCCAAGCTTCCTTTTTTGGTCAGTTTTTCGGTTAGAGGCTATGTTAACTACGATAAGCATGTAGCTAA
GTATTTAAAAATCTTGCCAGACCAGTTTTCTTATCAGGCAATAGAAGATGTTGTCAAAGCAGATGCTGA
AAAAAATACTTCCAATAATGAGATGGGAATGGAAAATTATTTCTATAATGAGCAGATCAAGAAGGATTT
GAAGAAATTAAAGGATTCTCAGAAAAGCTTTACCTATCTCAAGTCGCCAGAGTATAATGNNNTGCGATT
GGTTTTTAACACAGTTTTTCTAAATCTAAGGTAAACCCGATTTTTTATCATTCCACCTGTTAATAAAAAATG
GATGNACTATGCTGGTCTACGAGAGGATATGTACCAACAAACGGTGCAGAAGATTGCTACCAGTTAGA
AAGTCAAGGTTTTTACCAATATAGCAGATTTTTTCTAAGGACGGCGGGGAGCCTTTCTTTATGAAGGACAC
CATTCACCTTGGTTGGTTGGGTTGGTTGGCTTTTGGACAAGGCAGTTGATCCTTTCTATCCAATCCAC
ACCAGCTCCGACTTACCATCTGAATGAGCGCTTTTTTCAGCAAAGATTGGGCGACTTATGATGGAGATGT
CAAAGAA

SP038 amino acid (SEQ ID NO:58)

TEMHHNLGAEKRSVAVATTIDSFKERSQKVRALSDPNVRVFPFFGSSEWLRFDGAHSAVLAEKYNRSYRP
YLLGQGGAAASLNQYFGMQMLPQLENKQVVYVISPOWF SKNGYDPAAFQQYFNGDQLTSFLKHQSGDQA
SYAATRLLQQFPNVAMKDLVQKLASKEELSTADNEMI ELLARFNERQASFFGQFSVRGVVNYDKHVAK
YLKILPDQFSYQAIEDVVKADAENKTSNNEMGMENYFYNEQIKKDLKKLKDSQKSFTYLLKSPYXNLQL
VLTQFSKSKVNPFIPIPPVNKKWMXYAGLREDMYQQTVQKIRYQLESQGFTNIADFSKDGGEPPFMKDT
IHLGWLGLAFDKAVDPFLSNPTPAPTYHLNERFFSKDWATYDGDVKE

SP039 nucleotide (SEQ ID NO:59)

GGTTTTGAGAAAGTATTTGCAGGGGGCCCTGATTGAGTCGATTGAGCAAGTGGAAAATGACCGTATTGT
GGAAATTACAGTTTCCAATAAAAACGAGATTGGAGACCATATCCAGGCTACCTTGATTATCGAAATTAT
GGGGAAACACAGTAATATTCTACTGGTCGATAAAAGCAGTCATAAAATCCTCGAAGTTATCAAACACGT
CGGCTTTTTCACAAAATAGCTACCGCACCTTACTTCCAGGATCGACCTATATCGCTCCGCCAAGTACAAA
ATCTCTCAATCCTTTTACTATCAAGGATGAAAAGCTCTTTGAAATCCTGCAACCCCAAGAACTAACAGC
AAAAAATCTTCAAAGCCTCTTTCAAGGTCTGGGACGCGATACGGCAAATGAATTGGAAAGGATACTGGT
TAGTGAAAAACTTTCCGCTTTCCGAAATTTTTTCAATCAAGAAACCAAGCCATGCTTGACTGAGACTTC
CTTCAGTCCAGTTCCTTTTGCAATCAGGTGGGAGAGCCTTTTGCAAATCTTTCTGATTTGTTGGACAC
CTACTATAAGGATAAGGCTGAGCGCGACCGCTCAAACAGCAGGCCAGTGAAGTGAATTCGTCGTGTTGA
AAATGAACCTTCAGAAAAACCGACACAAACTCAAAAAACAGGAAAAAGAGTTACTGGCGACAGACAACGC
TGAAGAATTTCGTCAAAAAGGAGAATTGCTGACAACCTTCTCCACCAAGTGCCTAACGACCAAGACCA
GGTTATCTTAGCAACTACTATACCAACCAACCTATCATGATTGCGCTTGATAAGGCTCTGACTCCCAA
CCAGAATGCCAACGCTATTTTAAACGGTATCAGAAACTCAAAGAAGCTGTCAAATCTGACTGATTT
GATTGAAGAAACCAAGCCACTATTCTCTATCTGGAAAGTGTAGAAACCGTCCCAACCAAGCTGGACT
GGAAGAAATCGCTGAAATCCGTGAAGAATTGATTCAAACAGGTTTTATCCGCAGAAGACAACGGGAGAA
AATCCAGAAACGCAAAAAACTAGAACAAATATCTAGCAAGCGATGGCAAACCATCATCTATGTCCGACG
AAACAATCTTCAAATGAGGAATTGACCTTTAAATGGCCCGCAAGGAGGAACCTTGGTTCCATGCTAA
GGACATTCCTGGAAGCCATGTTGTCATCTCAGGAAATCTTGACCCATCTGATGCAGTCAAGACAGACGC

AGCAGAGTTAGCTGCCTACTTCTCTCAAGGGCGCCTGTGCAATCTGGTGCAGGTAGATATGATTGAAGT
CAAAAACTCAATAAACCAACTGGTGGAAAACCCGGCTTTGTCACTTACACAGGACAAAAGACCCTCCG
CGTCACACCAGACTCCAAAAAATTGCATCCATGAAAAATCC

SP039 amino acid (SEQ ID NO:60)

VLRKYLGALIESIEQVENDRIVEITVSNKNEIGDHIQATLIIIEIMGKHSNILLVDKSSHKILEVIKHV
GFSQNSYRLLPGSTYIAPPSTKSLNPFTIKDEKLFEILQTQELTAKNLQSLFQGLGRDTANELERILV
SEKLSAFRNFFNQETKPCLTETSFSPVPFANQVGEFFANLSDLLDTYYKDKAERDRVKQQASELIRRV
NELQKNRHKLKKQEKELLATDNAEEFRQKGELLTTFHQPNDQDQVILDNYITNQPIMIALDKALTPN
QNAQRYFKRYQKLKEAVKYLTDLIEETKATILYLESVETVLNQAGLEEIAEIREELIQTGFIRRRQREK
IQKRKKLEQYLASDGKTIIVYVGRNNLQNEELTFKMARKEELWFHAKDIPGSHVVISGNLDPDAVKTD
AELAAVFSQGRLSNLVQVDMIEVKLNKPTGGKPGFVITYTGQKTLRVTPDSKKIASMKKS

SP040 nucleotide (SEQ ID NO:61)

GACAACATTTACTATCCATACAGTAGAGTCAGCACCAGCAGAAGTGAAAGAAATTCTTGAAACAGTAGA
AAAAGACAACAATGGCTATATTCCCAACCTAATCGGTCTCTTGCCCAATGCCCCGACTGTTTAGAAGC
CTACCAAATTGTCTCATCTATCCACCGTCGCAACAGCCTGACACCCGTTGAGCGTGAAGTGGTGCAAAT
CACGGCAGCCGTGACCAATGGTTGTGCCTTCTGTGTCGAGGTACACAGCCTTTTCCATCAAACAAAT
CCAGATGAATGATGACTTGATTCAAGCTCTTCGCAATCGTACTCCAATTGAAACAGATCCTAAATTGGA
TACCCTAGCTAAGTTTACCTTGGCAGTTATCAATACCAAGGGTCGTGTAGGAGATGAAGCCTTGCTGA
GTTTTTAGAAGCTGGCTACACTCAACAAAAATGCCTTGGATGTGGTTTTTGGTGTGAGCCTAGCAATCCT
CTGTAACATATGCCAACAACCTTAGCTAATACCAATTAATCCAGAATTGCAACCTTATGCC

SP040 amino acid (SEQ ID NO:62)

TTFTIHTVESAPAEVKEILETVEKDNNGYIPNLIGLLANAPTIVLEAYQIVSSIHRNSLTPVEREVVQI
TAAVTNGCAFCVAGHTAFSIKQIQMNDLILQALRNRTPIETDPKLDLAKFTLAVINTKGRVGDALSE
FLEAGYTQQNALDVVFGVSLAILCNYANNLANTPINPELQPYA

SP041 nucleotide (SEQ ID NO:63)

GGCTAAGGAAAGAGTGGATGTACTAGCTTATAAACAGGGGTGTTTGAAACGAGAGAGCAGGCCAAGCG
AGGTGTGATGGCTGGCCTAGTCGTAGCAGTCCCTTAATGGAGAACGGTTTGACAAGCCAGGAGAGAAAAT
TCCAGATGACACCGAATTAACAACTCAAGGGGGAGAACTCAAGTATGTACAGCCGTGGTGGTTTGAACT
GGAAAAGGCCCTTGCAAGTCTTTGATTTGTGCGGTGGATGGCGCGACTACGATTGATATCGGGGCCTCTAC
TGGAGGTTTTTACCGATGTCTACAGAACTAGTGCCAAAGTTGGTCTTTGCAGTGCATGTTGGTACCAA
TCAGTTGGCTTGGAATTACGCCAAGACCCAGAGTTGTGAGCATGGAGCATGTTCAATTTCCGCTATGC
TGAAAAGACTGATTTTCGAGCAGGAGCCGAGCTTTGCCAGTATTGATGTGAGTTTCATTTCCCTTAGTCT
GATTTTGCCAGCCTTGACCCGTGTCTTGGCTGATCAAGGTCAGGTGGTAGCACTTGTCAAACCTCAGTT
TGAGGCAGGACGTGAGCAGATTGGGAAAAATGGAATTATTCGAGATGCTAAGGTTTCATCAGAATGTCCCT
TGAATCTGTAACAGCTATGGCAGTAGAGGTAGGTTTTTCAGTCCTTGGCTTGGACTTTTCTCCCATCCA
AGGTGGACATGGAATATTGAATTTTACGCTATTTGAAAAAAGAAAAGTCAGCAAGCAATCAGATTCT
TGCTGAGATTAAAGAAGCAGTAGAGAGGGCGCATAGTCAATTTAAAAATGAA

SP041 amino acid (SEQ ID NO:64)

AKERVDVLAYKQGLFETREQAKRGVMAGLVVAVLNGERFDKPGEKIPDDTELKLKGEKLYVSRGGLKL
EKALQVFDLSVDGATTIDIGASTGGFTDVMLQNSAKLVFAVDVGTNQLAWKLRQDPRVVSMEQFNFRYA
EKTDFEQEPSFASIDVSFISLSLILPALHRVLADQGVVALVKPQFEAGREQIGKNGIIRDAKVHQNVL
ESVTAMAVEVGFVSLGLDFSPIQGGHGNIEFLAYLKKEKSASNQILAEIKEAVERAHSQFKNE

SP042 nucleotide (SEQ ID NO:65)

TTGTTCCCTATGAACCTGGTCGTACCAAGCTGGTCAGGTTAAGAAAGAGTCTAATCGAGTTTCTTATAT
AGATGGTGATCAGGCTGGTCAAAAGGCAGAACTTGACACCAGATGAAGTCAGTAAGAGGGAGGGGAT
CAACGCCGAACAAATNGTNATCAAGATTACGATTCAAGGTTATGTGACCTCTCATGGAGACCATATCA
TTACTATAATGGCAAGGTTTCTTATGATGCCATCATCAGTGAAGAGCTCCTCATGAAAGATCCGAATTA
TCAGTTGAAGGATTCAGACATTGTCAATGAAATCAAGGGTGGTTATGTCAATTAAGGTAAACGGTAAATA
CTATGNTACCTTAAGGATGCAGCTCATGCGGATAATATTCGGACAAAAGAAGAGATTAAACGTCAGAA
GCAGGAACGCAGTCATAATCATAACTCAAGAGCAGATAATGCTGTTGCTGCAGCCAGAGCCCAAGGACG
TTATACAACGGATGATGGGTATATCTTCAATGCATCTGATATCATTGAGGACACGGGTGATGCTTATAT
CGTTCCTCACGGCGACCATTACCATTACATTCCTAAGAATGAGTTATCAGCTAGCGAGTTAGCTGCTGC

AGAAGCCTATTGGAATGGGAAGCAGGGATCTCGTCCTTCTTCAAGTTCTAGTTATAATGCAAATCCAGC
TCAACCAAGATTGTCAGAGAACCACAATCTGACTGTCACTCCAATTATCATCAAAATCAAGGGGAAAA
CATTTCAAGCCTTTTACGTGAATTGTATGCTAAACCCCTTATCAGAACGCCATGTGGAATCTGATGGCCT
TATTTTCGACCCAGCGCAAATCACAAGTGAACCGCCAGAGGTGTAGCTGTCCCTCATGGTAACCATTA
CCACTTTATCCCTTATGAACAAATGTCTGAATTGGAACCAAGTCCACAATCGACTCCGGAACCTAG
TCGTTCAAACCATTTGGGTACCAGATTCAAGACCAGAACAACCAAGTCCACAATCGACTCCGGAACCTAG
TCCAAGTCCGCAACCTGCACCAAATCCTCAACCAGCTCCAAGCAATCCAATTGATGAGAAATTTGGTCAA
AGAAGCTGTTTCGAAAAGTAGGCGATGGTTATGTCTTTGAGGAGAATGGAGTTTCTCGTTATATCCCAGC
CAAGGATCTTTTACGAGAAACAGCAGCAGGCATTGATAGCAAATGGCCAAGCAGGAAAGTTTATCTCA
TAAGCTAGGAGCTAAGAAAAGTACCTCCCATCTAGTGATCGAGAATTTTACAATAAGGCTTATGACTT
ACTAGCAAGAATTCACCAAGATTTACTTGATAATAAAGGTGACAAGTTGATTTTGAGGCTTTTGATAA
CCTGTTGGAACGACTCAAGGATGTCNCAAGTGATAAAGTCAAGTTAGTGGANGATATCTTGCCTTCTT
AGCTCCGATTCGTTCATCCAGAACGTTTAGGAAAACCAAATGCGCAAATTACCTACACTGATGATGAGAT
TCAAGTAGCCAAGTTGGCAGGCAAGTACACAACAGAAGACGGTTATATCTTTGATCCTCGTGATATAAC
CAGTGATGAGGGGGATGCCTATGTAACTCCACATATGACCCATAGCCACTGGATTAAGAAAAGATAGTTT
GTCTGAAGCTGAGAGAGCGGCAGCCAGGCTTATGTAAAGAGAAAGGTTTGACCCCTCCTTCGACAGA
CCATCAGGATTACAGAAATACTGAGGCAAAAGGAGCAGAAGCTATCTACAACCGCGTGAAAGCAGCTAA
GAAGGTGCCACTTGATCGTATGCCTTACAATCTTCAATATACTGTAGAAGTCAAAACCGGTAGTTTAAT
CATACCTCATTATGACCATTACCATAACATCAAATTTGAGTGGTTTGACGAAGGCCTTTATGAGGCACC
TAAGGGGTATACTCTTGAGGATCTTTTGGCGACTGTCAAGTACTATGTGCAACATCCAAACGAACGTC
GCATTCAGATAATGGTTTTTGGTAAACGCTAGCGACCATGTTCAAAGAAACAAAAATGGTCAAGCTGATAC
CAATCAAACGAAAAACCAAGCGAGGAGAAACCTCAGACAGAAAAACCTGAGGAAGAAACCCCTCGAGA
AGAGAAACCGCAAAGCGAGAAACAGAGTCTCCAAACCAACAGAGGAACAGAGAATCACCAGAGGA
ATCAGAAGAACCTCAGGTCGAGACTGAAAAGGTGAAGAAAACTGAGAGAGGCTGAAGATTTACTTGG
AAAAATCCAGGAT

SP042 amino acid (SEQ ID NO:66)

CSYELGRHQAGQVKKESNRVSYIDGDQAGQKAENLTPDEVSKREGINAEQXVIKITDQGYVTSHGDHYH
YYNGKVPYDAII SEELLMKDPNYQLKDS DIVNEIKGGYVIKVNGKYYVYLKDAAHADNIRTKEEIKRQK
QERSHNHNSRADNAVAAARAQGRYTTDDGYIFNASDII EDTGDAYIVPHGDHYHYI PKNELSASELAAA
EAYWNGKQGSRPSSSSSYNANPAQPR LSENHNLTVTPTYHQNGENISSLLRELYAKPLSERHVESDGL
IFDPAQITSRTARGVAVPHGNHYHFIPYEQMSELEKRIARI I PLRYRSNHWVPDSRPEQSPQSTPEPS
PSPQPAPNPQPAPSNPID EKL VKEAVRVKVG DGYVFEENGVSRYI PAKDL SAETAAGIDSKLAKQESLSH
KLGA KKTDL PSSDREFY NKAYD LLARI HQD LLDNKG RQVDFEALDNLLERLKD VXS DKVKLVXDIL AFL
APIRHPERL GKPN AQITYT DDEIQVAKLAGKYTTEDGYIFDPRDITSDEGDAYVTPHMTSHWIKKDSL
SEAERAAAQAYAKEKGLTPPSTDHQDSGNTEAKGAEAIYNRVKA AKKVPLDRMPYNLQYTV EVKNGSLI
IPHYDHYHNIKFEWFDEGLYEAPKGYTLEDLLATVKYVVEHPNERPHSDNGFGNASDHVQRNKNQADT
NQTEKPSEEEKPQTEKPEEETPREEK PQSEKPESPKPTEEPEESPEESEEPQVETEKVEEKLREAEDLLG
KIQD

SP043 nucleotide (SEQ ID NO:67)

TTATAAGGGTGAATTAGAAAAAGGATACCAATTTGATGGTTGGGAAATTTCTGGTTTCAAGGTAAAAA
AGACGCTGGCTATGTTATTAATCTATCAAAAGATACCTTTATAAAACCTGTATTCAAGAAAAATAGAGGA
GAAAAAGGAGGAAGAAAATAAACCTACTTTTGATGTATCGAAAAAGAAAGATAACCCACAAGTAAACCA
TAGTCAATTAAATGAAAGTCACAGAAAAAGAGGATTTACAAAGAGAAGAGCATTACAAAAATCTGATTC
AACTAAGGATGTTACAGCTACAGTTCTTGATAAAAAACAATATCAGTAGTAAATCAACTACTAACAATCC
TAATAAG

SP043 amino acid (SEQ ID NO:68)

YKGELEKGYQFDGWEISGFEGKKDAGYVINLSKDTFIKPVFKKIEEKKEENKPTFDVSKKKDNPQVNH
SQLNESHKEDLQREEHSQKSDSTKDV TATVLDKNNISSKSTNNPNK

SP044 nucleotide (SEQ ID NO:69)

GAATGTTCAAGGCTCAAGAAAGTTCAAGGAAATAAAATCCACTTTATCAATGTTCAAGAAAGGTGGCAGTGA
TGCGATTATTCTTGAAAGCAATGGACATTTTGCCATGGTGGATACAGGAGAAGATTATGATTTCCCAGA
TGGAAGTGATTCTCGCTATCCATGGAGAGAAGGAATTGAAACGTCTTATAAGCATGTTCTAACAGACCG
TGTCTTTTCGTCGTTTGAAGGAATTGGGTGTCCAAAACTTGATTTTATTTTGGTGACCCATACCCACAG
TGATCATATTGGAATGTTGATGAATTACTGTCTACCTATCCAGTTGACCGAGTCTATCTTAAGAAATA

TAGTGATAGTCGTATTACTAATTCTGAACGCTCTATGGGATAATCTGTATGGCTATGATAAGGTTTTACA
 GACTGCTGCAGAAAAAGGTGTTTCAGTTATTCAAATATCACACAAGGGGATGCTCATTTTTAGTTTGG
 GGACATGGATATTCAGCTCTATAATTATGAAAATGAACTGATTCATCGGGTGAATTAAAGAAAATTTG
 GGATGACAATTCCAATTCCTTGATTAGCGTGGTGAAAGTCAATGGCAAGAAAATTTACCTTGGGGGCGA
 TTTAGATAATGTTTCATGGAGCAGAAGACAAGTATGGTCTCTCATTGGAAAAGTTGATTGATGAAGTT
 TAATCATCACCATGATACCAACAAATCAAATACCAAGGATTTTATTAAAAATTTGAGTCCGAGTTTGAT
 TGTTCAAACCTTCGGATAGTCTACCTTGGAAAATGGTGGTGGATAGTGAAGTATGTTAATTGGCTCAAAGA
 ACGAGGAATTGAGAGAATCAACGCAGCCAGCAAAGACTATGATGCAACAGTTTTTGGATATTCGAAAAGA
 CGGTTTTGTCAATATTTCAACATCCTACAAGCCGATTCCAAGTTTTCAAGCTGGTTGGCATAAGAGTGC
 ATATGGGAAGTGGTGGTATCAAGCGCCTGATTCTACAGGAGAGTATGCTGTGCGTTGGAATGAAATCGA
 AGGTGAATGGTATTACTTTAACCACGCGGTATCTTGTACAGAATCAATGGAAAAATGGAACAATCA
 TTGGTTCTATTTGACAGACTCTGGTGCTTCTGCTAAAAATTTGGAAGAAAATCGCTGGAATCTGGTATTA
 TTTTAACAAAGAAAACCAGATGGAAATTGGTTGGATTCAAGATAAAGAGCAGTGGTATTTTGGATGT
 TGATGGTTCTATGAAGACAGGATGGCTTCAATATATGGGGCAATGGTATTACTTTGCTCCATCAGGGGA
 A

SP044 amino acid (SEQ ID NO:70)

NVQAQESSGNKIHFINVQEGGSDAIILESNGHFAMVDTGEDYDFPDGSDSRYPWREGIETSYKHVLTDR
 VFRRLKELGVQKLDLILVTHTHSDHIGNVDELLSTYFVDRVYLKKYSDSRITNSERLWDNLGYDKVLQ
 TAAEKGVSVIQNITQGDHAFQFGMDIQLYNYENETDSSGELKKIWDDNSNLSISVVKVNGKKIYLGDD
 LDNVHGAEDKYGPLIGKVDLMKFNHHDHTNKSNTKDFIKNLSPLIVQTSDSLPPWKNVGDSEYVNWLKE
 RGIERINAASKDYDATVDFDIRKDGfVNISTSYKPIPSFQAGWHKSAYGNWWYQAPDSTGEYAVGWNEIE
 GEWYYFNQTGILLQNWKKWNHWFYLTDSGASAKNWKKIAGIWIYYFNKENQMEIGWIQDKEQWYYLDV
 DGSMKTGWLQYMQWYFAPSGE

SP045 nucleotide (SEQ ID NO:71)

CTTGGGTGTAACCCATATCCAGCTCCTTCCAGTCTTGTCTTACTACTTTGTCAATGAATTGAAAAACCA
 TGAACGCTTGTCTGACTACGCTTCAAGCAACAGCAACTACAACCTGGGGATATGACCCCAAACTACTT
 CTCCTTGACTGGTATGTACTCAAGCGATCCTAAGAATCCAGAAAAACGAATCGCAGAATTTAAAAACCT
 CATCAACGAAATCCACAAACGTTGGTATGGGAGCTATCCTAGATGTCGTTTATAACCACACAGCCAAAGT
 CGATCTCTTTGAAGATTTGGAACCAAACTACTACCCTTTATGGATGCCGATGGCACACCTCGAACTAG
 CTTTGGTGGTGGACGCTTGGGGACAACCCACCATATGACCAAACGGCTCCTAATTGACTCTATCAAATA
 CCTAGTTGATACCTACAAAGTGGATGGCTTCCGTTTCGATATGATGGGAGACCATGACGCCGCTTCTAT
 CGAAGAAGCTTACAAGGCTGCACGCGCCCTCAACTGCAAACTCATCATGCTTGGTGAAGGTTGGAGAAC
 CTATGCCGGTGTGAAAACATGCCCTACTAAAGCTGCTGACCAAGATTGGATGAAACATACCGATCTGT
 CGCTGTCTTTTCAGATGACATCCGTAACACCTCAAATCTGGTTATCCAAACGAAGGTCAACCTGCCTT
 TATCACAGGTGGCAAGCGTGATGTCAACACCATCTTTAAAAATCTCATTGCTCAACCAACTAATTTGA
 AGCTGACAGCCCTGGAGATGTATCCAATACATCGCAGCCCATGATAACTTGACCTCTTTGACATCAT
 TGCCAGTCTATCAAAAAGACCCAAGCAAGGCTGAGAAGTATGCTGAAATCCACCGTCGTTTACGACT
 TGGAAATCTCATGGTCTTGACAGCTCAAGGAAGTCCATTTATCCACTCCGGTCAGGAATATGGACGTAC
 TAAACAATTCCGTGACCCAGCCTACAAGACTCCAGTAGCAGAGGATAAGGTTCCAAACAAATCTCACTT
 GTTGCGTGATAAGGACGGCAACCCATTTGACTATCCTTACTTCATCCATGACTCTTACGATCTTAGTGA
 TGCAGTCAACAAGTTTGACTGGACTAAGGCTACAGATGGTAAAGCTTATCCTGAAAATGTCAAGAGCCG
 TGACTATATGAAAGGTTTGATTGCCCTTCGTCAATCTACAGATGCCTTCCGACTTAAGAGTCTTCAAGA
 TATCAAAGACCGTGTCCACCTCATCACTGTCCCAGGCCAAAATGGTGTGGAAAAAGAGGATGTAGTGAT
 TGGCTACCAAATCACTGCTCCAAACGGCGATATCTACGCAGTCTTTGTCAATGCGGATGAAAAAGCTCG
 CGAATTTAATTTGGGAAGTGCCTTTGCACATCTAAGAAATGCGGAAGTTTGGCAGATGAAAAACCAAGC
 AGGACCAGTCGGAATTGCCAACCCGAAAGGACTTGAATGGACTGAAAAAGGCTTGAAATGAATGCCCT
 TACAGCTACTGTTCTTCGAGTCTCTCAAAATGGAAGTACCATGAGTCAACTGCAGAAGAGAAACCAGA
 CTCAACCCCTTCCAAGCCTGAACATCAAAATGAAGCTTCTCACCCTGCACATCAAGACCCAGCTCCAGA
 AGCTAGACCTGATTCTACTAAACCAGATGCCAAAGTAGCTGATGCGGAAAATAAACCTAGCCAAGCTAC
 AGCTGATTACAAAGCTGAACAACCAGCACAGAAGCACAAGCATCATCTGTAAAAGAAGCGGTTTCGAAA
 CGAATCGGTAGAAAACCTTAGCAAGGAAAATATACCTGCAACCCAGATAAACAAGCTGAA

SP045 nucleotide (SEQ ID NO:72)

LGVTHIQLLPVLSYYFVNELKNHERLSYASSNSNYNWGYDPQNYFSLTGMYSDDPKNPEKRIAEFKNL
 INEIHKRGMGAILDVVYNHTAKVDLFEDLEPNYYHFMDADGTPRTSFGGGRLGTHHMTKRLLIDSIKY
 LVDTYKVDGFRFDMGDHDAASIEEAYKAARALNPNIIMLGEGWRTYAGDENMPTKAADQDWMKHTDTV

AVFSDDIRNNLKSYPNEGQPAFITGGKRDVNTIFKNLIAQPTNFEADSPGDV IQYIAAHDNLTFLDII
AQSIKKDPSKAENYAEIHRRLRLGNLMVLTAQGTFFIHSGQEYGR TKQFRDPAYKTPVAEDKVPNKSHL
LRDKDGNPFDPYFPIHDSYDSSDAVNKF DWT KATD GKAYPENVKSRDYMKG LIALRQSTDAFRLKSLQD
IKDRVHLITVPGQNGVEKEDVVIGYQITAPNGDIYAVFVNADEKAREFN LGTAF AHLRNAEVLADENQA
GPVGIANPKGLEWTEKGLKLNALTATVLRVSQNGTSHESTAEEKPDSTPSKPEHQNEASHPAHQDPAPE
ARPDSTKPD AKVADAENKPSQATADSQA EQPAQEAQASSVKEAVRNESVENSSKENIPATPDKQAE

SP046 nucleotide (SEQ ID NO:73)

TAGTGATGGTACTTGGCAAGGAAAACAGTATCTGAAAGAAGATGGCAGTCAAGCAGCAAATGAGTGGGT
TTTNGATACTCATTATCAATCTTGGTTCCTATATAAAAGCAGATGCTAACTATGCTGAAAATGAATGGCT
AAAGCAAGGTGACGACTATTTTTACCTCAAATCTGGTGGCTATATGGCCAAATCAGAATGGGTAGAAGA
CAAGGGAGCCTTTTATTATCTTGACCAAGATGGAAAGATGAAAAGAAATGCTTGGGTAGGAACCTCCTA
TGTTGGTGCAACAGGTGCCAAAAGTAATAGAAGACTGGGTCTATGATTCTCAATACGATGCTTGGTTTTA
TATCAAAGCAGATGGACAGCACGCAGAGAAAGAATGGCTCCAAATTAAAGGGAAGGACTATTATTTCAA
ATCCGGTGGTTATCTACTGACAAGTCAGTGGATTAATCAAGCTTATGTGAATGCTAGTGGTGCCAAAGT
ACAGCAAGGTTGGCTTTTTGACAAACAATACCAATCTTGGTTTTACATCAAAGAAAATGGAACTATGC
TGATAAAGAATGGATTTTCGAGAATGGTCACTATTATTATCTAAAATCCGGTGGCTACATGGCAGCCAA
TGAATGGATTGGGATAAGGAATCTTGGTTTTATCTCAAATTTGATGGGAAAATGGCTGAAAAAGAATG
GGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGGTGGTTACATGACAGCCAATGAATG
GATTTGGGATAAGGAATCTTGGTTTTACCTCAAATCTGATGGGAAAATAGCTGAAAAAGAATGGGTCTA
CGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCTGGTGGCTACATGGCGAAAATGAGACAGTAGA
TGGTTATCAGCTTGAAGCGATGGTAAAATGGCTTGGAGGAAAAACTACAAATGAAAATGCTGCTTACTA
TCAAGTAGTGCCTGTTACAGCCAATGTTTATGATTCAGATGGTGAAAAGCTTTCTATATATCGCAAGG
TAGTGTCGTATGGCTAGATAAGGATAGAAAAAGTGATGACAAGCGCTTGGCTATTACTATTTCTGGTTT
GTCAGGCTATATGAAAACAGAAGATTTACAAGCGCTAGATGCTAGTAAGGACTTTATCCCTTATTATGA
GAGTGATGGCCACCGTTTTTATCACTATGTGGCTCAGAATGCTAGTATCCCAGTAGCTTCTCATCTTTC
TGATATGGAAGTAGGCAAGAAATATTATTCGGCAGATGGCCTGCATTTTGATGGTTTTAAGCTTGAGAA
TCCCTTCCTTTTCAAAGATTTAACAGAGGCTACAAACTACAGTGCTGAAGAATTGGATAAGGTATTTAG
TTTGCTAAACATTAACAATAGCCTTTTGGAGAACAAGGGCGCTACTTTTAAGGAAGCCGAAGAACATTA
CCATATCAATGCTCTTTATCTCCTTGCCCATAGTGCCCTAGAAAAGTAACTGGGGAAGAAAGTAAAAATTGC
CAAAGATAAGAATAATTTCTTTGGCATTAACAGCCTATGATACGACCCCTTACCTTTCTGCTAAGACATT
TGATGATGTGGATAAGGGAATTTTAGGTGCAACCAAGTGGATTAAGGAAAATTATATCGATAGGGGAAG
AACTTTCCTTGGAACAAGGCTTCTGGTATGAATGTGGAATATGCTTCAGACCCCTTATTGGGGCGAAAA
AATTGCTAGTGTGATGATGAAAATCAATGAGAAGCTAGGTGGCAAAGAT

SP046 amino acid (SEQ ID NO:74)

SDGTWQ GKQYLKEDGSQAANEVVDTHYQSWFYIKADANYAENEWLKQGDDYFY LKSGGYMAKSEWVED
KGAFYYLDQDGKMKRNAWVGTSYVGATGAKVIEDWVYDSQYDAWFYIKADGQHAKEKWLQIKGKDYYFK
SGGYLLTSQWINQAYVNASGAKVQGWLF DKQYQSWFYIKENGNYADKEWIFENGHYYYLKSGGYMAAN
EWIWDKESWFY LKF DGKMAEK EWVYDSHSQAWY YFKSGGYMTANEWIWDKESWFY LKSDGKIAEK EWVY
DSHSQAWY YFKSGGYMAKNETVDGYQLGSDGKWLGGKTTNENAAYYQVVPVTANVYDS DGEKLSYISQ
SVVWLDKDRKSDDKRLAITISGLSGYMKTEDLQALDASKDFIPYYESDGH RFYHYVAQNASIPVASHLS
DMEVGKKYYSADGLHFDGFKLENPFLFKDLTEATNYSAEELDKVFSLLNINNSLLENKGATFKEAEEHY
HINALYLLAHSALSNWGRSKI AKDKNNFFGITAYDTPYLSAKTFDDVDKGILGATKWIKENYIDRGR
TFLGNKASGMNVEYASDPYWGEKIASVMMKINEKLGGKD

SP048 nucleotide (SEQ ID NO:75)

TGGGATTCAATATGTCAGAGATGATACTAGAGATAAAGAAGAGGGAATAGAGTATGATGACGCTGACAA
TGGGGATATTATTGTAAAAGTAGCGACTAAACCTAAGGTAGTAACCAAGAAAATTTCAAGTACGCGAAT
TCGTTATGAAAAAGATGAAACAAAAGACCGTAGTGAATCCTGTTACAATTGATGGAGAGGATGGCTA
TGTAACTAGCACAAGGACCTACGATGTTAATCCAGAGACTGGTTATGTTACCGAACAGGTTACTGTTGA
TAGAAAAGAAGCCACGGATACAGTTATCAAAGTTCCAGCTAAAAGCAAGGTTGAAGAAGTCTCTGTTCC
ATTTGCTACTAAATATGAAGCAGACAATGACCTTTCTGCAGGACAGGACGAAGAGATTACTCTAGGAAA
GAATGGGAAAACAGTTACAACGATAA CTTATAATGTAGATGGAAAGAGTGGACAAGTAACCTGAGAGTAC
TTTAAGTCAAAAAAAGACTCTCAAACAAGAGTTGTTAAAAAAGaACCArkCCCCAAGTTCTTGTCCA
AGAAATTCCAATCGAAACAGAATATCTCGATGGCCCACTCTTGATAAAaGTCAAGAAAGTAGAAGAAGT
AGGAGAAATTGGTAAATTACTCTTACTACAATCTATACTGGTAGATGAACGTGATGGAACAATTGAAGA
AACTACTTCTCGTCAAATTACTAAAGAGATGGTAAAAGACGTATAAGGAGAGGGACGAGAGAACCTGA

AAAAGTTGTTGTTCTCCTGAGCAATCATCTATTCCTTCGTATCCTGTATCTGTTACATCTAACCAAGGAAC
AGATGTAGCAGTAGAACCAGCTAAAGCAGTTGCTCCAACAACAGACTGGAAACAAGAAAATGGTATGTG
GTATTTTATAATACTAGTGGTTCCATGGCAACAGGTTGGGTACAAGTTAATAGTTCATGGTACTACCT
CAACAGCAACGGTTCTATGAAAGTCAATCAATGGTTCCAAGTTGGTGGTAAATGGTATTATGTAAATAC
ATCGGGTGAGTTAGCGGTCAATACAAGTATAGATGGCTATAGAGTCAATGATAATGGTGAATGGGTGCG
T

SP048 amino acid (SEQ ID NO:76)

GIQYVRDDTRDKEEGIEYDDADNGDIIVKVATKPKVVTKKISSSTRIRYKDETKDRSENPVITIDGEDGY
VTTTTRTYDVNPETGYVTEQVTVDRKEATDTVIKVPKSKVEEVLVPFATKYEADNDLSAGQEQEITLGK
NGKTVTTTITYNVDGKSGQVTESTLSQKKDSQTRVVKKRTXPQVLVQEIPIETEYLDGPTLDKSQEVVEV
GEIGKLLLLQSILVDERDGTIEETTSRQITKEMVKRRIIRRGTREPEKVVVPEQSSIPSPVSVTSNQG
DVAVEPAKAVAPTTDWKQENGMWYFYNTDGSMATGWVQVNSSWYYLNSNGSMKVNQWFQVGGKWWYVNT
SGELAVNTSIDGYRVNDNGEVR

SP049 nucleotide (SEQ ID NO:77)

GGATAATAGAGAAGCATTAACCTTTATGACGGGTGAAAATTTTATCTCCAACATTATCTAGGAGC
ACATAGGGAAGAACTAAATGGAGAGCATGGCTATACCTTCCGTGTTTGGGCACCTAATGCTCAGGCTGT
TCATTGGTTGGTGTATTTACCAACTGGATTGAAAATCAGATTCCAATGGTAAGAAATGATTTTGGGGT
CTGGGAAGTCTTTACCAATATGGCTCAAGAAGGCATATTTACAAATATCATGTACACGCTCAAAATGG
TCATCAACTGATGAAGATTGACCTTTTGCCTGTCAGGTATGAGGCTCGTCCAGGAACAGGGGCAATCGT
AACAGAGCTTCTGAGAAGAAATGGAAGGATGGACTTTGGCTGGCAGCAAGAAAACGTTGGGGCTTTGA
AGAGCGTCTGTCAATATTTATGAAGTTCACGCTGGATCATGGAAGAAATCTGATGGCAGTCCTTA
TAGTTTGGCCAGCTCAAGGATGAAGTCAATTCCTTATCTCGTTGAAATGAAGTATACTCATATTGAGTT
TATGCCCTTGATGTCCCATCCTTTGGGCTTGAGTTGGGGGTATCAGCTTATGGGTTACTTCGCTTTAGA
GCATGCTTATGGCCGACCAGAGGAGTTTCAAGATTTTGTC

SP049 amino acid (SEQ ID NO:78)

DNREALKTFMTGENFYLLQHYLGAAHREELNGEHGYTFRVWAPNAQAVHLVGDFTNWIENQIPMVRNDFGV
WEVFTNMAQEGHIYKYHVTRQNGHQLMKIDPFAVRYEARPGTGAIVTELPEKKWKDGLWLARRKRWGF
ERPVIYEVHAGSWKRNSDGSPYSFAQLKDELIPLYVEMNYTHIEFMPLMSHPLGLSWGYYQLMGYFALE
HAYGRPEEFQDFV

SP050 nucleotide (SEQ ID NO:79)

AGATTTTGTGCGAGGAGTGTACATACCCATAATATTGGGGTTATTGTGGACTGGGTACCAGNTCACTTTAC
CATCAACGATGATGCCTTAGCCTATTATGATGGGACACCGACTTTTGAATACCAAGACCATAATAAGGC
TCATAACCATGGTTGGGGTGCCCTTAATTTTACCTTGGAAAAATGAAGTCCAGTCCTTCTTAATTTCT
TTGCATTAAGCATTGGATTGATGTCTATCATTTGGATGGTATTCGTGTGGATGCTGTTAGCAACATGCT
CTATTTGGACTATGATGATGCTCCATGGACACCTAATAAAGATGGCGGAAATCTCAACTATGAAGGTTA
TTATTTCTTTCAGCGCTTGAATGAGGTATTAAGTTAGAATATCCAGATGTGATGATGATTGCAGAAGA
AAGTTCGTCTGCGATCAAGATTACGGGAATGAAAGAGATTGGTGGTCTAGGATTTGACTACAAATGGAA
CATGGGCTGGATGAATGATATCCTCCGTTTCTACGAAGAAGATCCGATCTATCGTAAATATGACTTTAA
CCTGGTGAATTTTCACTTTATGTATGTTNCAAGGAGAATTATCTCTTGCCATTCTCGCACGATGAAGT
GGTTCATGGCAAGAAGAGTATGATGCATAAGATGTGGGGAGATCGTTACAATCAATTCGCAGGCTTGCG
CAATCTCTATACGTACCAATTTGTACCCCTGGTAAGAAATTGCTCTTCATGGGTAGCGAATACGGTCA
ATTCCTAGAATGGAATCTGAAGAACAGTTGGAATGGTCTAACCTAGAAGACCCAATGAATGCTAAGAT
GAAGTATTTTCGCTTCTCAGCTAAACAGTTTACAAAGATCATCGCTGTCTGTGGGAAATTGATACCAG
CTATGATGGTATTGAAATCATTGATGCGGATAATCGAGACCAGAGTGTCTTTCTTTTATTCGTAAGGG
TAAAAAGGGA

SP050 amino acid (SEQ ID NO:80)

DFVEECHTHNIGVIVDWVPXHFTINDDALAYYDGTPTFEYQDHNKAHNHGWGALNFDLGKNEVQSFLIS
CIKHWDIVYHLDGIRVDAVSNMMLYLDYDDAPWTPNKDGGNLNYEGYYFLQRLNEVIKLEYPDVMMIAEE
SSSAIKITGMKEIGGLGFDYKWNMGWMNDILRFYEEDPIYRKYDFNLVTF SFMYVXKENYLLPF SHDEV
VHGKKSMMHKMWGDRYNQFAGLRNLYTYQICHPGKKLLFMGSEYGQFLEWKSEEQLEWSNLEDPMNAKM
KYFASQLNQFYKDHRLWEIDTSYDGIEIIDADNRDQSVLSFIRKGGK

SP051 nucleotide (SEQ ID NO:81)

60

ATCTGTAGTTTATGCGGATGAAACACTTATTACTCATACTGCTGAGAAACCTAAAGAGGAAAAAATGAT
AGTAGAAGAAAAGGCTGATAAAGCTTTGGAAACTAAAAATATAGTTGAAAGGACAGAACAAAGTGAACC
TAGTTCAACTGAGGCTATTGCATCTGAGNAGAAAGAAGATGAAGCCGTAACCTCAAAAAGAGGAAAAAGT
GTCTGCTAAACCGGAAGAAAAAGCTCCAAGGATAGAATCACAAGCTTCAAATCAAGAAAAACCGCTCAA
GGAAGATGCTAAAGCTGTAACAAATGAAGAAGTGAATCAAATGATTGAAGACAGGAAAGTGGATTTTAA
TCAAAATTGGTACTTTAAACTCAATGCAAATTTAAGGAAGCCATTAAACCTGATGCAGACGTATCTAC
GTGGAAAAAATTAGATTTACCGTATGACTGGAGTATCTTTAACGATTTTCGATCATGAATCTCCTGCACA
AAATGAAGGTGGACAGCTCAACGGTGGGGAAGCTTGGTATCGCAAGACTTTCAAAGTATGAAAAAGA
CCTCAAGAAAAATGTTTCGCTTACTTTTGTATGGCGTCTACATGGATTCTCAAGTTTATGTCAATGGTCA
GTTAGTGGGGCATTATCCAAATGGTTATAACCAGTTCTCATATGATATCACCAAATACCTTCAAAAAGA
TGGTCGTGAGAATGTGATTGCTGTCCATGCAGTCAACAAACAGCCAAGTAGCCGTTGGTATTCAGGAAG
TGGTATCTATCGTGATGTGACTTTACAAGTGACAGATAAGGTGCATGTTGAGAAAAATGGGACAACCTAT
TTTAACACCAAACCTTGAAGAACAACAACATGGCAAGGTTGAAACTCATGTGACCAGCAAAATCGTCAA
TACGGACGACAAAGACCATGAACTTGTAGCCGAATATCAAATCGTTGAACGAGGTGGTCAATGCTGTAAC
AGGCTTAGTTTCGTACAGCGAGTCGTACCTTAAAAGCACATGAATCAACAAGCCTAGATGCGATTTTAGA
AGTTGAAAGACCAAACTCTGGACTGTTTTAAATGACAAACCTGCCTTGTACGAATTGATTACGCGTGT
TTACCGTGACGGTCAATTGGTTGATGCTAAGAAGGATTTGTTTGGTTACCGTTACTATCACTGGACTCC
AAATGAAGGTTTCTCTTTGAATGGTGAACGTATTAAATTCATGGAGTATCCTTGCACCACGACCATTG
GGCGCTTGGAGCAGAAGAAAACTATAAAGCAGAATATCGCCGTCTCAAACAAATGAAGGAGATGGGAGT
TAACCTCCATCCGTACAACCCACAACCTGCTAGTGAGCAAAACCTTGCAAATCGCAGCAGAATAGGTTT
ACTCGTTTCAGGAAGAGGCCTTTGATACGTGGTATGGTGGCAAGAAACCTTATGACTATGGACGTTTCTT
TGAAAAAGATGCCACTCACCCAGAAGCTCGAAAAGGTGAAAAATGGTCTGATTTTGACCTACGTACCAT
GGTCGAAAGAGGCAAAAACAACCTGCTATCTTCATGTGGTCAATTGGTAATGAAATAGGTGAAGCTAA
TGGTGATGCCCACTCTTTAGCAACTGTTAAACGTTTGGTTAAGGTTATCAAGGATGTTGATAAGACTCG
CTATGTTACCATGGGAGCAGATAAATCCGTTTCGGTAATGGTAGCGGAGGGCATGAGAAAATTGCTGA
TGAACCTCGATGCTGTTGGATTTAACTATCTGAAGATAATTACAAAGCCCTTAGAGCTAAGCATCCAAA
ATGGTTGATTTATGGATCAGAAACATCTTCAGCTACCCGTACACGTGGAAGTTACTATCGCCCTGAACG
TGAATTGAAACATAGCAATGGACCTGAGCGTAATTATGAACAGTCAGATTATGGAAATGATCGTGTGGG
TTGGGGGAAAACAGCAACCGCTTCATGGACTTTTGACCGTGACAACGCTGGCTATGCTGGACAGTTTAT
CTGGACAGGTACGGACTATATTGGTGAACCTACACCATGGCACAACCAAAATCAAACCTCCTGTTAAGAG
CTCTTACTTTGGTATCGTAGATACAGCCGGCATTCAAAACATGACTTCTATCTCTACCAAAGC

SP051 amino acid (SEQ ID NO:82)

SVVYADETLITHTAEKPKEEKMIVEEKADKALETKNIVERTEQSEPSSTEAIASEXKEDEAVTPKEEKV
SAKPEEKAPRIESQASNQEKPLKEDAKAVTNEEVNQMIEDRKVDFNQNWYFKLNANSKEAIKPDADVST
WKKLDLPYDWSIFNDFDHESPAQNEGGQLNGGEAWYRKTFKLDEKDLKKNVRLTFDGVYMDSQVYVNGQ
LVGHYPNGYNQFSYDITKYLQKDGRENVIAVHAVNKQPSSRWYSGSGIYRDVTLQVTDKVHVEKNGTTI
LTPKLEEQQHGKVVETHVTSKIVNTDDKDHELVAEYQIVERGGHAVTGLVRTASRTLKAHESTSLDAILE
VERPKLWTVLNDKPALYELITRVYRDGQLVDAKKDLFGYRYHWTNPNEGFSLNGERIKFHGVSLLHHDHG
ALGAEENYKAEYRRLKQMKEMGVNSIRTHNPASEQTLQIAAELGLLVQEEAFDTWYGGKPYDYGRFF
EKDATHPEARKGEKWSDFDLRTMVERGKNNPAIFMWSIGNEIGEANGDAHSLATVKRLVKVIKDVKTR
YVTMGADKFRFGNGSGGHEKIADELDAVGFNYSYEDNYKALRAKHPKWLIIYGETSSATRTRGSYYRPER
ELKHSNGPERNYEQSDYGNDRVGWGKTATASWTFDRDNAGYAGQFIWTGTDYIGEPTPWHNQNPVKS
SYFGIVDTAGIPKHDFLYQS

SP052 nucleotide (SEQ ID NO:83)

TTACTTTGGTATCGTAGATACAGCCGGCATTCAAAACATGACTTCTATCTCTACCAAAGCCAATGGGT
TTCTGTTAAGAAGAAACCGATGGTACACCTTCTTCTCACTGGAACCTGGGAAAAACAAAGAAATTAGCATC
CAAAGTAGCTGACTCAGAAGGTAAGATTCCAGTTCTGTGCTTATTTCGAATGCTTCTAGTGTAGAATTGTT
CTTGAATGGAAAAATCTCTTGGTCTTAAGACTTTCAATAAAAAACAAACCAGCGATGGGCGGACTTACCA
AGAAGGTGCAAAATGCTAATGAACCTTATCTTGAATGGAAAGTTGCCTATCAACCAGGTACCTTGGAGC
AATTGCTCGTGAATGTAATCTGGCAAGGAAATGCTCGAGATAAGATTACGACTGCTGGTAAGCCAGCGGC
AGTTTCGTCTTATTAAGGAAGACCATGCGATTGTCAGCAGATGGAAAGACTTGACTTACATCTACTATGA
AATTGTTGACAGCCAGGGGAATGTGGTTCCAACCTGCTAATAATCTGGTTTCGCTTCCAATTGCAATGGCCA
AGGTCAACTGGTCGGTGTAGATAACCGGAGAACAAGCCAGCCGTGAACGCTATAAGGCGCAAGCAGATGG
TTCTTGGATTTCGTAAAGCATTTAATGGTAAAGGTGTTGCCATTGTCAAATCAACTGAACAAGCAGGGAA
ATTCACCTGACTGCCCACTCTGATCTCTTGAATCGAACCAAGTCACTGTCTTTACTGGTAAGAAAGA
AGGACAAGAGAAGACTGTTTTGGGGACAGAAGTGCCAAAAGTACAGACCATTATTGGAGAGGCACCTGA

67

"E80F9680" 1083 60E0F

AATGCCTACCACTGTTCCGTTTGTATACAGTGATGGTAGCCGTGCAGAACGTCCTGTAACCTGGTCTTC
AGTAGATGTGAGCAAGCCTGGTATTGTAACGGTGAAAGGTATGGCTGACGGACGAGAAGTAGAAGCTCG
TGTAAGAAGTGATTGCTCTTAAATCAGAGCTACCAGTTGTGAAACGTATTGCTCCAAATACTGACTTGAA
TTCTGTAGACAAATCTGTTTCCCTATGTTTTGATTGATGGAAGTGTGAAGAGTATGAAGTGGACAAGTG
GGAGATTGCCGAAGAAGATAAAGCTAAGTTAGCAATTCCAGGTTCTCGTATTCAAGCGACCGGTATTT
AGAAGGTCAACCAATTCATGCAACCCCTGTGGTAGAAGAAGGCAATCCTGCGGCACCTGCAGTACCAAC
TGTAACGGTTGGTGGTGAGGCAGTAACAGGTCTTACTAGTCAAAAACCAATGCAATACCGCACTCTTGC
TTATGGAGCTAAGTTGCCAGAAGTCACAGCAAGTGCTAAAAATGCAGCTGTTACAGTTCTTCAAGCAAG
CGCAGCAAACGGCATGCGTGCGAGCATCTTTATTACGCCTAAAGATGGTGGCCCTCTTCAAACCTATGC
AATTCAATTCCCTTGAAGAAGCGCCAAAAATTGCTCACTTGAGCTTGCAAGTGGAAAAAGCTGACAGTCT
CAAAGAAGACCAAACTGTCAAATTGTCGGTTCGAGCTCACTATCAAGATGGAACGCAAGCTGTATTACC
AGCTGATAAAGTAACCTTCTCTACAAGTGGTGAAGGGGAAGTCGCAATTCGTAAAGGAATGCTTGAGTT
GCATAAGCCAGGAGCAGTCACTCTGAACGCTGAATATGAGGGAGCTAAAGACCAAGTTGAACCTCACTAT
CCAAGCCAATACTGAGAAGAAGATTGCGCAATCCATCCGTCCTGTAAATGTAGTGACAGATTTGCATCA
GGAACCAAGTCTTCCAGCAACAGTAACAGTTGAGTATGACAAAGGTTTCCCTAAAACTCATAAAGTCAC
TTGGCAAGCTATTCCGAAAGAAAACTAGACTCCTATCAAACATTTGAAGTACTAGGTAAAGTTGAAGG
AATTGACCTTGAAAGCGGTGCAAAAGTCTCTGTAGAAGGTATCGTTTCAGTTGAAGAAGTCAGTGTGAC
AATCCAATCGCAGAAGCACCACAATTACCAGAAAGTGTTCCGACATATGATTCAAATGGTCAAGTTTC
ATCAGCTAAGGTTGCATGGGATGCGATTTCGTCCAGAGCAATACGCTAAGGAAGGTGCTTTTACAGTTAA
TGGTCGCTTAGAAGGTACGCAATTAACA

SP052 amino acid (SEQ ID NO:84)

YFGIVDTAGIPKHDFYLYQSQWVSVKKKPMVHLLPHWNWENKELASKVADSEGKIPVRAYSNAASSVELF
LNGKSLGLKTFNKKQTS DGRTYQEGANANELYLEWKVAYQPGTLEAIARDESGKEIARDKITTAGKPAA
VRLIKEDHAIADGKDLTYIYYEIVDSQGNVPTANNLVRFLHGOQQLVGVNDNEQASRERYKAQADG
SWIRKAFNGKGVAIVKSTEQAGKFTLTAHSDLLKSNQVTVFTGKKEGQEKTVLGTEVPKVQTIIGEAPE
MPTTVPFVYS DGSRAERPVTWSSVDVSKPGIVTVKGMADGREVEARVEVIALKSELPPVKRIAPNTDLN
SVDKSVSYVLIDGSVEEYEVDKWEIAEEDKAKLAI PGSRIQATGYLEGQPIHATLVVEEKNPAAPAVPT
VTVGGEAVTGLTSQKPMQYRTLAYGAKLPEVTASAKNAAVTVLQASAANGMRASIFIQPKDGGPLQTYA
IQFLEEAPKIAHLSLQVEKADSLKEDQTVKLSVRAHYQDGTQAVLPADKVTFSTSGEGEVAIRKGMLEL
HKPGAVTLNAEYEGAKDQVELTIQANTEKKIAQSIRPVNVVTDLHQEPSLPATVTVEYDKGFPKTHKVT
WQAIPEKLDYSYQTFEVLGKVEGIDLEARAKVSVVEGIVSVVEEVSVTTPIAEAPQLPESVRTYDSNGHVS
SAKVAWDAIRPEQYAKEGVFTVNGRLEGTQLT

SP053 nucleotide (SEQ ID NO:85)

AGCTAAGGTTGCATGGGATGCGATTCTGTCAGAGCAATACGCTAAGGAAGGTGTCTTTACAGTTAATGG
TCGCTTAGAAGGTACGCAATTAACAACCTAACTTTCATGTTTCGCGTATCTGCTCAAACTGAGCAAGGTGC
AAACATTTCTGACCAATGGACCGGTTCAAGATTGCCACTTGCCCTTTGCTTCAGACTCAAAATCCAAGCGA
CCCAGTTTCAAATGTAAATGACAAGCTCATTTCTTACAATAACCAACCAGCCAATCGTTGGACAACTG
GAATCGTACTAATCCAGAAGCTTCAGTCGGTGTCTGTTTGGAGATTACAGTATCTTGAGCAAACGCTC
CGTTGATAATCTAAGTGTGCGATTCCATGAAGACCATGGAGTTGGTGTACCGAAGTCTTATGTGATTGA
GTATTATGTTGGTAAGACTGTCCCAACAGCTCTTAAAAACCCCTAGTTTGTGTTGGTAATGAGGACCATGT
CTTTAATGATTCTGCCAACTGGAAACCAGTTACTAATCTAAAAGCCCCTGCTCAACTCAAGGCTGGAGA
AATGAACCACTTTAGCTTTGATAAAGTTGAAACCTATGCTGTTTCGTATTTCGCATGGTTAAAGCAGATAA
CAAGCGTGGAACGTCTATCACAGAGGTACAAATCTTTGCGAAACAAGTTGCGGCAGCCAAGCAAGGACA
AACAAGAATCCAAGTTGACGGCAAAAGACTTAGCAAACTTCAACCCTGATTTGACAGACTACTACCTTGA
GTCTGTAGATGGAAGGTTCCGGCAGTCACAGCAAGTGTTAGCAACAATGGTCTCGCTACCGTCGTTCC
AAGCGTTTCGTGAAGGTGAGCCAGTTTCGTGTATCGCGAAAGCTGAAAATGGCGACATCTTAGGAGAATA
CCGTCTGCACTTCACTAAGGATAAGAGCTTACTTTCTCATAAAACAGTTGCTGCGGTTAAACAAGCTCG
CTTGCTACAAGTAGGTCAAGCACTTGAATTGCCGACTAAGGTTCCAGTTTACTTCACAGGTAAAGACGG
CTACGAACAAAAGACCTGACAGTTGAATGGGAAGAAGTTCCAGCGGAAAATCTGACAAAAGCAGGTCA
ATTTACTGTTTCGAGGCCGTGTCCTTGGTAGTAACCTTGTGCTGAGATCACTGTACGAGTGACAGACAA
ACTTGGTGAGACTCTTTAGATAACCTTAACCTATGATGAAAACAGTAACAGGCCCTTTGCTTCAGCAAC
CAATGATATTGACAAAACTCTCATGACCGCGTTGACTATCTCAATGACGGAGATCAATTCAGAAAATCG
TCGTTGGACAACTGGTCACCAACACCATCTTCTAATCCAGAAGTATCAGCGGGTGTGATTTTCCGTGA
AAATGGTAAGATTGTAGAACGGACTGTTACACAAGGAAAAGTTCAAGTTCTTTGAGATAGTGGTACGGA
TGCACCATCTAACTCGTTTTAGAACGCTATGTCGGTCCAGAGTTTGAAGTGCCAACCTACTATTCAA
CTACCAAGCCTACGACGCAGACCATCCATTCAACAATCCAGAAAATTGGGAAGCTGTTCTTATCGTGC

GGATAAAGACATTGCAGCTGGTGTATGAAATCAACGTAACATTTAAAGCTATCAAAGCCAAAGCTATGAG
ATGGCGTATGGAGCGTAAAGCAGATAAGAGCGGTGTTGCGATGATTGAGATGACCTTCCTTGCACCAAG
TGAATTGCCTCAAGAAAGCACTCAATCAAAGATTCTTGTAGATGGAAAAGAACTTGCTGATTTTCGCTGA
AAATCGTCAAGACTATCAAATTACCTATAAAGGTCAACGGCCAAAAGTCTCAGTTGAAGAAAACAATCA
AGTAGCTTCAACTGTGGTAGATAGTGGAGAAGATAGCTTTCCAGTACTTGTTCGCCTCGTTTCAGAAAG
TGGAAAACAAGTCAAGGAATACCGTATCCACTTGACTAAGGAAAAACCAGTTTCTGAGAAGACAGTTGC
TGCTGTACAAGAAGATCTTCCAAAAATCGAATTTGTTGAAAAAGATTTGGCATAACAAGACAGTTGAGAA
AAAAGATTCAACACTGTATCTAGGTGAAACTCGTGTAGAACAAGAAGGAAAAGTTGAAAAAGAACGTAT
CTTTACAGCGATTAATCCTGATGGAAGTAAGGAAGAAAAACTCCGTGAAGTGGTAGAAGTCCGACAGA
CCGCATCGTCTTGGTTGGAACCAACCAGTAGCTCAAGAAGCTAAAAAACCAAGTGTGAGAAAAAGC
AGATACAAAACCAATTGATTCAAGTGAAGCTAGTCAAACATAATAAGCCAG

SP053 amino acid (SEQ ID NO:86)

AKVAWDAIRPEQYAKEGVFTVNGRLEGTQLTKLHVRVSAQTEQGANISDQWTGSELPLAFASDSNPSPD
PVSNNVDKLISSNNQPANRWTNWNRTNPEASVGLFGDSGILSKRSVDNLSVGFHEDHGVGVPKSYVIE
YYVGKTVPAPKNPSFVGNEDHVFND SANWKPVNTNLKAPQLKAGEMNHFSFDKVETVAVRIRMVKADN
KRGTSITEVQIFAKQVAAAKQGQTRIQVDGKDLANFNPDLDYYLESDVGKVPVAVTASVSNNGLATVVP
SVREGEPVRVIAKAENGDI LGEYRLHFTKDKSLLSHKPVA AVKQARLLQVGQALELPTKVPVYFTGKDG
YETKDLTVEWEEVPAENLTAKAQFTVRGRVLSNLVAEITVRVTDKLGELSDNPNYDENSNNQAFASAT
NDIDKNSHDRVDYLDNDGDHSENRRWTNWSPTSSNPEVSAGVIFRENGKIVERTVTQGVQFFADSGTD
APSKLVRLERYVGPEFEVPTYYSNYQAYDADHPFNPNPENWEAVPYRADKDIAAGDEINVTFFKAIKAKAMR
WRMERKADKSGVAMIEMTFLAPSELPQESTQSKILVDGKELADFAENRQDYQITYKGQRPKVSVENNO
VASTVVDSDGSDSFPVLVRLVSESGKQVKEYRIHLTKEKPVSEKTVA AVQEDLPKIEFVEKDLAYKTVEK
KDSTLYLGETRVEQEGKVGKERIFTAINPDGSKEEKLREVEVPTDRIVLVGTPVAQEAKKPQVSEKA
DTKPIDSSEASQTNKAQ

SP054 nucleotide (SEQ ID NO:87)

CTATCACTATGTAAATAAAGAGATTATTTTCAACAAGACTAAAGATTTAATTCAGACAGGAAAGCCTGA
CAGGAATGAAGTTGTATATGGTTTGGTGTATCAAAAAGATCAGTTGCCTCAAACAGGGACAGAA

SP054 amino acid (SEQ ID NO:88)

YHYVNKEIISQEAKDLIQTKPDRNEVVYGLVYQKDQLPQTGTE

SP055 nucleotide (SEQ ID NO:89)

TGAGACTCCTCAATCAATAACAAATCAGGAGCAAGCTAGGACAGAAAACCAAGTAGTAGAGACAGAGGA
AGCTCCAAAAGAAGAAGCACCTAAAACAGAAGAAAGTCCAAAGGAAGAACCAAAATCGGAGGTAAAACC
TACTGACGACACCCTTCTCTAAAGTAGAAGAGGGGAAAGAAGATTACAGCAGAACCAGCTCCAGTTGAAGA
AGTAGGTGGAGAAGTTGAGTCAAAAACCAGAGGAAAAAGTAGCAGTTAAGCCAGAAAAGTCAACCATCAGA
CAAACCAGCTGAGGAATCAAAAGTTGAACAAGCAGGTGAACCAGTCGCGCCAAGAGAAGACGAAAAGGC
ACCAGTCGAGCCAGAAAAGCAACCAGAAGCTCCTGAAGAAGAGAAGGCTGTAGAGGAAAACACCGAAACA
AGAAGAGTCAACTCCAGATACCAAGGCTGAAGAACTGTAGAACCAAAAGAGGAGACTGTTAATCAATC
TATTGAACAACCAAAAGTTGAAACGCCTGCTGTAGAAAAACAACAGAACCAACAGAGGAACCAAAAGT
TGAACAAGCAGGTGAACCAGTCGCGCCAAGAGAAGACGAACAGGCACCAACGGCACCAGTTGAGCCAGA
AAAGCAACCAGAAGTTCTGGAAGAAGAGAAGGCTGTAGAGGAAACACCGAAACCAGAAGATAAAATAAA
GGGTATTGGTACTAAAGAACCAGTTGATAAAAGTGAGTTAAATAATCAAATTGATAAAGCTAGTTCAGT
TTCTCCTACTGATTAT

SP055 amino acid (SEQ ID NO:90)

ETPQSITNQEQARTENQVVETEEAPKEEAPKTEESPKEEPKSEVKPTDDTLPKVEEGKEDSAEPAPVEE
VGGEVESKPEEKVAVKPESQPSDKPAEESKVEQAGEPVAPREDEKAPVEPEKQPEAPEEEKAVEETPKQ
EESTPDTKAEETVEPKEETVNQSI EQPKVETPAVEKQTEPTEEPKVEQAGEPVAPREDEQAPTAPVEPE
KQPEVPEEEKAVEETPKPEDKIKGIGTKEPVDKSELNNQIDKASSVSPTDY

SP056 nucleotide (SEQ ID NO:91)

GGATGCTCAAGAAACTGCGGGAGTTCACTATAAATATGTGGCAGATTACAGAGCTATCATCAGAAGAAAA
GAAGCAGCTTGTCTATGATATTCCGACATACGTGGAGAATGATGATGAACTTATTATCTTGTTTATAA
GTTAAATTCTCAAATCAACTGGCGGAATTGCCAAATACTGGAAGCAAGAATGAGAGGCAA

SP056 amino acid (SEQ ID NO:92)

DAQETAGVHYKYVADSELSSEEKKQLVYDIPTYVENDDETYLVYKLNSQNQLAELPNTGSKNERQ

SP057 nucleotide (SEQ ID NO:93)

CGACAAAGGTGAGACTGAGGTTCAACCAGAGTCGCCAGATACTGTGGTAAGTGATAAAGGTGAACCAGA
GCAGGTAGCACCGCTTCCAGAATATAAGGGTAATATTGAGCAAGTAAACCTGAAACTCCGGTTGAGAA
GACCAAAGAACAAGGTCCAGAAAAAACTGAAGAAGTTCAGTAAAACCAACAGAAACACCAGTAAA
TCCAAATGAAGGTACTACAGAAGGAACCTCAATTCAAGAAGCAGAAAAATCCAGTTCAACCTGCAGAAGA
ATCAACAACGAATTCAGAGAAAAGTATCACCAGATACATCTAGCAAAAATACTGGGGAAGTGTCCAGTAA
TCCTAGTGATTGCGACAACCTCAGTTGGAGAATCAAATAAACCAGAACATAATGACTCTAAAAATGAAAA
TTCAGAAAAAACTGTAGAAGAAGTTCAGTAAATCCAAATGAAGGCACAGTAGAAGGTACCTCAAATCA
AGAAACAGAAAAACCAGTTCAACCTGCAGAAGAAACACAAACAACTCTGGGAAAAATAGCTAACGAAAA
TACTGGAGAAGTATCCAATAAACCTAGTGATTCAAACCACCAGTTGAAGAATCAAAATCAACCAGAAAA
AAACGGAACCTGCAACAAAAACCAGAAAAATTCAGGTAATACAACATCAGAGAATGGACAAAACAGAACCA
ACCATCAAACGGAAATTCAACTGAGGATGTTTCAACCGAATCAAACACATCCAATTCAAATGGAAACGA
AGAAATTAAACAAGAAAATGAACTAGACCCGTATAAAAAGGTAGAAGAACCAGAGAAAAACACTTGAATT
AAGAAAT

SP057 amino acid (SEQ ID NO:94)

DKGETEVQPESPDTVVSDKGEPEQVAPLPEYKGNIEQVKPETPVEKTKEQGPEKTEEVVVKPTEETPVN
PNEGTTGTSIQEAENPVQPAEESTTNSEKVSPTSSKNTGEVSSNPSDSTTSVGESNKPEHNDSKNEN
SEKTVEEVVNPNEGTVGTSNQETEKVPQPAEETQTN SGKIANENTGEVSNKPSDSKPPVEESNQPEK
NGTATKPENSGNTTSENGQTEPEPSNGNSTEDVSTESNTSNSNGNEEIKQENELDPDKKVEEPEKTLEL
RN

SP058 nucleotide (SEQ ID NO:95)

AAATCAATTGGTAGCACAAAGATCCAAAAGCACAAAGATAGCACTAAACTGACTGCTGAAAAATCAACTGT
TAAAGCACCTGCTCAAAGAGTAGATGTAAAAGATATAACTCATTTAACAGATGAAGAAAAAGTTAAGGT
TGCTATTTTACAAGCAAATGGTTTCAGCATTAGACGGAGCGACAATCAATGTAGCTGGAGATGGTACAGC
AACAATCACATTCCAGATGGTTTCAGTAGTGACGATTCCTAGGAAAAGATACAGTTCAACAATCTGCGAA
AGGTGAATCTGTAACCTCAAGAAGCTACACCAGAGTATAAGCTAGAAAATACACCAGGTGGAGATAAGGG
AGGCAATACTGGAAGCTCAGATGCTAATGCGAATGAAGGCGGTGGTAGCCAGGCGGGTGGATCAGCTCA
CACAGGTTCAAAAACCTCAGCTCAATCACAAGCTTCTAAGCAATTAGCTACTGAAAAAGAATCAGCTAA
AAATGCCATTGAAAAAGCAGCCAAGGACAAGCAGGATGAAATCAAAGGCGCACCGCTTCTGTATAAAGA
AAAAGCAGAACTTTTAGCAAGAGTGGAAGCAGAAAAACAAGCAGCTCTCAAAGAGATTGAAAATGCGAA
AACTATGGAAGATGTGAAGGAAGCAGAAACGATTGGAGTGCAAGCCATTGCCATGGTTACAGTTCCTAA
GAGACCAGTGGCTCCTAAT

SP058 amino acid (SEQ ID NO:96)

NQLVAQDPKAQDSTKLTAEKSTVKAPAQRVDVKDITHLTDEEKVKVAILQANGSALDGATINVAGDGTA
TITFPDGSVVTILGKDTVQSAKGESVTQEATPEYKLENTPGGDKGNTGSSDANANEGGGSQAGGSAH
TGSQNSAQSQASKQLATEKESAKNAIEKAAKDKQDEIKGAPLSDKEKAELLARVEAEKQALKEIENAK
TMEDVKEAETIGVQAIAMVTVPKRPVAPN

SP059 nucleotide (SEQ ID NO:97)

CAAACAGTCAGCTTCAGGAACGATTGAGGTTGATTTACAGAGAAAATGGCTCTGGGACACGGGGTGCCTT
CACAGAAATCACAGGGATTCTCAAAAAAGACGGTGATAAAAAAATTGACAACACTGCCAAAAACAGCTGT
GATTCAAAATAGTACAGAAGGTGTTCTCTCAGCAGTTCAAGGGAATGCTAATGCTATCGGCTACATCTC
CTTGGGATCTTTAACGAAATCTGTCAAGGCTTTAGAGATTGATGGTGTCAAGGCTAGTCGAGACACAGT
TTTAGATGGTGAATACCCTCTTCAACGTCCCTTCAACATTGTTTGGTCTTCTAATCTTTCCAAGCTAGG
TCAAGATTTTATCAGCTTTATCCACTCCAAACAAGGTCAACAAGTGGTCACAGATAATAAATTTATTGA
AGCTAAAACCGAAACCAAGGAATATACAAGCCAACACTTATCAGGCAAGTTGTCTGTTGTAGGTTCCAG
TTCAGTATCTCTTTAATGGAATAATAGCAGAAGCTTATAAAAAAGAAAAATCCAGAAGTTACGATTGA
TATTACCTCTAATGGGTCTTCAGCAGGTATTACCGCTGTTAAGGAGAAAACCGCTGATATTGGTATGGT
TTCTAGGGAATTAACCTCTGAAGAAGGTAAGAGTCTCACCATGATGCTATTGCTTTAGACGGTATTGC
TGTTGTGGTCAATAATGACAATAAGGCAAGCCAAGTCAGTATGGCTGAACCTGCAGACGTTTTTAGTGG
CAAATTAACCACCTGGGACAAGATTAAA

SP059 amino acid (SEQ ID NO:98)

KQSASGTIEVISRENGSGTRGAFTEITGILKKDGDKKIDNTAKTAVIQNSTEGVLSAVQGNANAIGYIS
LGS�TKSVKALEIDGVKASRDTVLDGEYPLQRPFNIVWSSNLKLGQDFISFIHKSQGGQVVTDNKFIE
AKTETTEYTSQHLSGKLSVVGSTSVSSLMEKLAAYKKENPEVTIDITSNGSSAGITAVKEKTADIGMV
SRELTPPEGKSLTHDAIALDGIADVNNNDNKASQVSMAELADVFSGLTTWWDKIK

SP060 nucleotide (SEQ ID NO:99)

ATTTCGATGATGCGGATGAAAAGATGACCCGTGATGAAATTGCCTATATGCTGACAAATAGTGAAGAAAC
ATTGGATGCTGATGAGATTGAGATGCTACAAGGTGTCTTTTCGCTCGATGAACTGATGGCACGAGAGGT
TATGGTTCCTCGAACGGATGCCTTTATGGTGGATATTCAGGATGATAGTCAAGCCATTATCCAAAGTAT
TTTAAAACAAAATTATTCTCGTATCCCGGTTTATGATGGGGATAAGGACAATGTAATTGGAATCATTCA
CACCAAGAGTCTCCTTAAGGCAGGCTTTGTGGACGGTTTTGACAATATTGTTTGAAGAGAATTTTACA
AGATCCACTTTTTGTACCTGAAACTATTTTTGTGGATGACTTGCTAAAAGAACTGCGAAATACCCAAAG
ACAAATG

SP060 amino acid (SEQ ID NO:100)

FDDADEKMTRDEIAYMLTNSEETLDADEIEMLQGVFSLDELMAREVMVPRTDAFMVDIQDDSQAI IQSI
LKQNYSRIPVYDGDKNVIGIIHTKSLKAGFVDGFDNIVWKRIQDPLFVPETIFVDDLLKELRNTQR
QM

SP062 nucleotide (SEQ ID NO:101)

GGAGAGTCGATCAAAAGTAGATGAAGCTGTGTCTAAGTTTGAAAAGGACTCATCTTCTTCGTCAAGTTC
AGACTCTTCCACTAAACCGGAAGCTTCAGATACAGCGAAGCCAAACAAGCCGACAGAACCAGGAGAAAA
GGTAGCAGAAGCTAAGAAGAAGGTTGAAGAAGCTGAGAAAAAGCCAAGGATCAAAAAGAAGAAGATCG
TCGTAACCTACCCAACCATTACTTACAAAACGCTTGAACCTGAAATTGCTGAGTCCGATGTGGAAGTTAA
AAAAGCGGAGCTTGAAGTAGTAAAAGTGAAAGCTAACGAACCTCGAGACGAGCAA

SP062 amino acid (SEQ ID NO:102)

ESRSKVDEAVSKFEKDS SSSSSSSSTKPEASDTAKPNKPTEPGEKVAEAKKKVEEA EKKAKDQKEEDR
RNYPTITYKTLELEIAESDVEVKAELELVKVKANEPRDEQ

SP063 nucleotide (SEQ ID NO:103)

ATGGACAACAGGAACTGGGACGAGGTTATATCTGGTAAGATTGACAAGTACAAAGATCCAGATATTCC
AACAGTTGAATCACAAGAAGTTACGTCAGACTCTAGTGATAAAGAAATAACGGTAAGGTATGACCGTTT
ATCAACACCAGAAAAACCAATCCCAACCAAAATCCAGAGCATCCAAGTGTTCGACACCAAAACCCAGA
ACTACCAAAATCAAGAGACTCCAACACCAGATAAACCAACTCCAGAACCCAGGTACTCCAAAACTGAAAC
TCCAGTGAATCCAGACCCAGAAAGTTCCGACTTATGAGACAGGTAAGAGAGAGGAATTGCCAAACACAGG
TACAGAAGCTAAT

SP063 amino acid (SEQ ID NO:104)

WTTGNWDEVISGKIDKYKDPDIPTVESQEVTS DSSDKEITVRYDRLSTPEKPI PQPNPEHPSVPTPNPE
LPNQETPTPKPTPEPGTPTKTETPVNPDPEVPTYETGKREELPNTGTEAN

SP064 nucleotide (SEQ ID NO:105)

CGATGGGCTCAATCCAACCCAGGTCAAGTCTTACCTGAAGAGACATCGGGAACGAAAGAGGGTGACTT
ATCAGAAAAACCAGGAGACACCGTTCTCACTCAAGCGAAACCTGAGGGCGTTACTGGAAATACGAATTC
ACTTCCGACACCTACAGAAAGAACTGAAGTGAGCGAGGAAACAAGCCCTTCTAGTCTGGATACACTTTT
TGAAAAAGATGAAGAAGCTCAAAAAAATCCAGAGCTAACAGATGTCTTAAAAGAACTGTAGATACAGC
TGATGTGGATGGGACACAAGCAAGTCCAGCAGAACTACTCCTGAACAAGTAAAAGGTGGAGTGAAAGA
AAATACAAAAGACAGCATCGATGTTCTCTGCTGCTTATCTTGAAAAAGCTGAAGGGAAAGGTCCCTTTCAC
TGCCGGTGTAACCAAGTAATTCTTATGAACTATTCGCTGGTGATGGTATGTTAACTCGTCTATTACT
AAAAGCTTTCGGATAATGCTCCTTGGTCTGACAATGGTACTGCTAAAAATCCTGCTTTACCTCCTCTTGA
AGGATTAACAAAAGGAAATACTTCTATGAAGTAGACTTAAATGGCAATACTGTTGGTAAACAAGGTCA
AGCTTTAATTGATCAACTTCGCGCTAATGGTACTCAAACCTATAAAGCTACTGTTAAAGTTTACGGAAA
TAAAGACGGTAAAGCTGACTTGACTAATCTAGTTGCTACTAAAAATGTAGACATCAACATCAATGGATT
AGTTGCTAAAGAAACAGTTCAAAAAGCCGTTGCAGACAACGTTAAAGACAGTATCGATGTTCCAGCAGC
CTACCTAGAAAAAGCCAAGGGTGAAGGTCCATTACAGCAGGTGTCAACCATGTGATTCCATACGAAC
CTTCGCAGGTGATGGCATGTTGACTCGTCTCTTGCTCAAGGCATCTGACAAGGCACCATGGTCAGATAA

CGGCGACGCTAAAAACCCAGCCCTATCTCCACTAGGCGAAAACGTGAAGACCAAAGGTCAATACTTCTA
TCAANTAGCCTTGGACGGAAATGTAGCTGGCAAAGAAAAACAAGCGCTCATTGACCAGTTCGAGCAAA
NGGTACTCAAACCTTACAGCGCTACAGTCAATGTCTATGGTAACAAAGACGGTAAACCAGACTTGGACAA
CATCGTAGCAACTAAAAAAGTCACTATTAACATAAACGGTTTAATTTCTAAAGAAAACAGTTCAAAAAGC
CGTTGCAGACAACGTTAANGACAGTATCGATGTTCCAGCAGCCTACCTAGAAAAAGCCAAGGGTGAAGG
TCCATTACAGCAGGTGTCAACCATGTGATTCATACGAACCTCTTCGAGGTGATGGTATGTTGACTCG
TCTCTTGCTCAAGGCATCTGACAAGGCACCATGGTCAGATAACGGNGACGCTAAAAACCCAGCNCATC
TCCACTAGGTGAAAACGTGAAGACCAAAGGTCAATACTTCTATCAANTAGCCTTGGACGGAAATGTAGC
TGGCAAAGAAAAACAAGCGCTCATTGACCAGTTCGAGCAAAACGGTACTCAAACCTTACAGCGCTACAGT
CAATGTCTATGGTAACAAAGACGGTAAACCAGACTTGGACAACATCGTAGCAACTAAAAAAGTCACTAT
TAAGATAAATGTTAAAGAAACATCAGACACAGCAAATGGTTTCATTATCACCTTCTAACTCTGGTTCTGG
CGTGACTCCGATGAATCACAATCATGCTACAGGTACTACAGATAGCATGCCTGCTGACACCATGACAAG
TTCTACCAACACGATGGCAGGTGAAAACATGGCTGCTTCTGCTAACAAGATGTCTGATACGATGATGTC
AGAGGATAAAGCTATG

SP064 amino acid (SEQ ID NO:106)

DGLNPTPGQVLPEETSGTKEGDLSEKPGDVLVTQAKPEGVGTGNTNSLPTPTERTVEVSEETSPSSLDTLF
EKDEEAQKNPELTDVLKETVDADVDGTQASPAETTPEQVKGKVNTKDSIDVPAAYLEKAEGKGPFT
AGVNQVIPEYLFAGDGMLTRLLKASDNAPWSDNKTAKNPALPPLLEGLTKGYFYEVLDNNGTVGKQGG
ALIDQLRANGTQTYKATVKVYGNKDGKADLTNLVATKNVDININGLVAKETVQKAVADNNGKDSIDVPA
YLEKAKGEGPFTAGVNHVIPEYLFAGDGMLTRLLKASDKAPWSDNGDAKNPALSPLGENVKTGQYFY
QXALDGNVAGKEKQALIDQFRAXGTQTYSATVNVYGNKDGKPDLDNIVATKKVTININGLISKETVQKA
VADNVXDSIDVPAAYLEKAKGEGPFTAGVNHVIPEYLFAGDGMLTRLLKASDKAPWSDNGDAKNPALS
PLGENVKTGQYFYQXALDGNVAGKEKQALIDQFRANGTQTYSATVNVYGNKDGKPDLDNIVATKKVTI
KINVKETSDTANGSLSPNSGSGVTPMNHNHATGTTDSMPADTMTSSNTMAGENMAASANKMSDTMMS
EDKAM

SP065 nucleotide (SEQ ID NO:107)

TTCCAATCAAAAACAGGCAGATGGTAAACTCAATATCGTGACAACCTTTTACCCTGTCTATGA_rTTTAC
CAAGCAAGTCGCAGGAGATACGGCTAATGTAGAAC_rTCCTAATCGGTGCTGGGACAGAACCTCATGAATA
CGAACCATCTGCCAAGGCAGTTGCCAAAAATCCAAGATGCAGATACCTTCGTTTATGAAAATGAAAACAT
GGAAACATGGGTACCTAAATTGCTAGATACCTTGGATAAGAAAAAAGTGA_rAACCATCAAGGCGACAGG
CGATATGTTGCTCTTGGCAGGTGGCGAGGAAGAAGAGGGAGACCATGACCATGGAGAAGAAGGTCATCA
CCATGAGTTTGACCCCCATGTTTGGTTATCACCAGTTCGTGCCAT_rAAACTAGTAGAGCACCATCCGCG
ACACTTGTGCAGCAGATTATCCTGATAAAAAAGAGACCTTTTGAGAAGAATGCAGCTGCCTATATCGAAAA
ATTGCAAGCCTTGGATAAGGCTTACGCAGAAGGTTTGTCTCAAGCAAAACAAAAGAGCTTTGTGACTCA
ACACGCAgCCTTTAACTaTCTTGCCTTGGACTATGGGACTC

SP065 amino acid (SEQ ID NO:108)

SNQKQADGKLNIVTTFFYPVYEFTKQVAGDTANVELLIGAGTEPHEYEPSAKAVAKIQDADTFVYENENM
ETWVPKLLD_rLDLKKVKTIKATGDMLLLPGGEEEEGDHHDHGEEGHHHEFDPHVWLS_rPVRAIKLVEHHP
HLSADY_rPKKETFEKNAAAYIEKLQALDKAYAEGLSQAKQKSFVTQHAAFNYLALDYGT

SP067 nucleotide (SEQ ID NO:109)

TATCACAGGATCGAACGGTAAGACAACCAACGACTATGATTGGGGAAGTTTTGACTGCTGCTGGCCA
ACATGGTCTTTTATCAGGGAATATCGGCTATCCAGCTAGTCAGGTTGCTCAAATAGCATCAGATAAGGA
CACGCTTGTTATGGAAC_rTTCTTCTTTCCAAC_rCATGGGTGTTCAAGAATTCATCCAGAGATTGCGGT
TATTACCAACCTCATGCCAACTCATATCGACTACCATGGGTCA_rTTTTCGGAATATGTAGCAGCCAAGTG
GAATATCCAGAACAAGATGACAGCAGCTGATTTCTTGTCTTGAAC_rTTAATCAAGACTTGGCAAAAGA
CTTGACTTCCAAGACAGAAGCCACTGTTGTACCATTTTCAACACTTGA_rAAAGGTTGATGGAGCTTATCT
GGAAGATGGTCAACTCTACTTCCGTGGTGAAGTAGTCATGGCAGCGAATGAAATCGGTGTTCCAGGTAG
CCACAATGTGGAAAAATGCCCTTGCAGACTATGTCTGTAGCCAAAGCTTCGTGATGTGGACAATCAAACCAT
CAAGGAAACTCTTTTACGCTTCCGTGGTGTC_rCAACACCGTCTCCAGT_rTTGTGGATGACATCAAGGGTGT
TAAATTC_rTATAACGACAGTAAATCAACTAATATCTTGGCTACTCA_rAAAAGCCTTGTCAAGGATTGACAA
CAGCAAGGTCGTCTTGATTGCAGGTGGTTTGGACCGTGGCAATGAGTTTGCAGAAATTGGTGCCAGACAT
TACTGGACTCAAGAAGATGGTCATCTGGGTCAATCTGCAGAACGTGTCAAACGGGCAGCAGACAAGGC
TGGTGTCTGCTTATGTGGAGGCGACAGATATTGCAGATGCGACCCGCAAGGCCTATGAGCTTGCAGCTCA

AGGAGATGTGGTTCTTCTTAGTCTGCCAATGCTAGCTGGGATATGTATGCTAACTTTGAAGTACGTGG
CGACCTCTTTATCGACACAGTAGCGGAGTTAAAAGAA

SP067 amino acid (SEQ ID NO:110)

GITGSNGKTTTTTMMIGEVLTAAGQHGLLSGNIGYPASQVAQIASDKDTLVMELSSFLMGVQEFHPEIA
VITNLMPTHIDYHGSFSEYVAAKWNIQNKMTAADFLVLNFNQDLAKDLTSKTEATVVPFSTLEKVDGAY
LEDGQLYFRGEVVMANEIGVPGSHNVENALATIAVAKLRDQDNQTIKETLSAFGGVKHRLQFVDDIKG
VKFYNDKSTNILATQKALSGFDNSKVVLIAAGGLDRGNEFDELVPDITGLKKMVLGQSAERVKRAADK
AGVAYVEATDIADATRKAYELATQGDVVLSPANASWDMYANFEVRGDLFIDTVAELKE

SP068 nucleotide (SEQ ID NO:111)

AAGTTCATCGAAGATGGTTGGGAAGTCCACTATATCGGGGACAAGTGTGGTATCGAACACCAAGAAATC
CTTAAGTCAGGTTTGGATGTACCTTCCATTCTATTGCGACTGGAAAATTGCGTCGCTATTCTCTTGG
CAAAATATGCTGGACGTCTTCAAAGTTGGTTGGGGAATTGTCCAATCGCTCTTTATCATGTTGCGACTG
CGTCCACAGACCCTTTTTTCAAAGGGGGGCTTTGTCTCAGTACCGCCTGTTATCGCTGCGCGTGTGTCA
GGAGTGCCTGTCTTTATTACGAATCTGACCTGTCTATGGGCTTGCCAATAAAATCGCCTATAAAATTT
CGGACTAAGATGTATTCAACCTTTGAACAAGCTTCGAGTTTGGCTAAGGTTGAGCATGTGGGAGCGG

SP068 amino acid (SEQ ID NO:112)

SSSKMVGKSTISGTSVVSNTKKSLSQVWMSPSILLRLENCVAISLGKICWTSSKLVGELSNRSLSCDCD
VHRPFFQRGALSQYRLSLRVCQECLSLFTNLTCWAWPIKSPINLRLRCIQPLNKLRLVWLRSLMWER

SP069 nucleotide (SEQ ID NO:113)

ATCGCTAGCTAGTGAAATGCAAGAAAGTACACGTAAATTCAAGGTTACTGCTGACCTAACAGATGCCGG
TGTTGGAACGATTGAAGTTCCTTTGAGCATTGAAGATTTACCCAATGGGCTGACCGCTGTGGCGACTCC
GCAAAAAATTACAGTCAAGATTGGTAAGAAGGCTCAGAAGGATAAGGTAAAGATTGTACCAGAGATTGA
CCCTAGTCAAATTGATAGTCGGGTACAAATTGAAAATGTCATGGTGTGAGATAAAGAAGTGCTATTAC
GAGTGACCAAGAGACATTGGATAGAATTGATAAGATTATCGCTGTTTTGCCAACTAGCGAACGTATAAC
AGGTAATTACAGTGGTTTCACTACCTTTGCAGGCAATCGACCGCAATGGTGTGTCTTACCGGCAGTTAT
CACTCCGTTTGATACAATAATGAAGGTGACTACAAAACAGTAGCACCAAGTTCAAGCACATCAAATTC
AAGTACAAGCAGTTCATCGGAGACATCTTCGTCAACGAAAGCAACTAGTTCAAAAACGAAT

SP069 amino acid (SEQ ID NO:114)

SLASEMQESTRKFKVTADLTDAGVGTIEVPLSIEDLPNGLTAVATPQKITVKIGKKAQKDKVKIVPEID
PSQIDSRVQIENVMVSDKEVSITSDQETLDRIDKIIAVLPTSERITGNYSVPLQAI DRNGVVLPAVI
TPFDTIMKVTTKPVAPSSSTNSSTSSSSETSSSTKATSSKTN

SP070 nucleotide (SEQ ID NO:115)

GCACCAGATGGGGCACAAGGTTCAAGGATCAGATGTTGAAAAGTACTACTTTACCCAACGCGGTCTTGA
GCAGGCAGGAATTACCATTCTTCTTTTGATGAAAAAATCTAGACGGTGATATGGAAATTATCGCTGG
AAATGCCTTTTCGTCCAGATAACAACGTCGAAATTGCCTATGCGGACCAAAATGGTATCAGCTACAAACG
TTACCATGAGTTTCTAGGTAGCTTTATGCGTGACTTTGTTAGCATGGGAGTAGCAGGAGCACATGGAAA
AACTTCAACGACAGGTATGTTGTCTCATGTCTTGTCTCACATTACAGATACCAGCTTCTTGATTGGAGA
TGGGACAGGTCGTGGTTTCGGCCAATGCCAAATATTTTGTCTTTGAATCTGACGAATATGAGCGTCACCT
CATGCCTTACCACCCAGAATACTCTATTATCACCAACATTGACTTTGACCATCCAGATTATTTACACAAG
TCTCGAGGATGTTTTTAATGCCTTTAACGACTATGCCAAACAAATCACCAAGGGTCTTTTTTGTCTATGG
TGAAGATGCTGAATTGCGTAAGATTACGTCTGATGCACCAATTTATTATTATGGTTTTGAAGCTGAAGG
CAATGACTTTGTAGCTAGTGATCTTCTTCGTTCAATAACTGGTTCAACCTTCACCGTTTCAATTTCCGTGG
ACAAAACCTTGGGGCAATTCCACATTCCAACCTTTGGTCGTCACAATATCATGAATGCGACAGCCGTTAT
TGGTCTTCTTTACACAGCAGGATTTGATTTGAACTTGGTGCGTGAGCACTTGAAAACATTTGCCGGTGT
TAAACGTCGTTTCACTGAGAAAATTGTCAATGATACAGTGATTATCGATGACTTTGCCCACCATCCAAC
AGAAATTATTGCGACCTTGGATGCGGCTCGTCAGAAATACCCAAGCAAGGAAATTGTAGCAGTCTTTCA
ACCGCATACCTTTACAAGAACCATTGCCTTGTGGAGCACTTTGCCCATGCTTTAAACCAAGCAGATGC
TGTTTTATCTAGCGCAAATTTATGGCTCGGCTCGTGAAGTAGATCATGGTGACGTTAAGGTAGAAGACCT
AGCCAACAAAATCAACAAAAACACCAAGTGATTACTGTTGAAAATGTTTCTCCACTCCTAGACCATGA
CAATGCTGTTTACGTCTTTATGGGAGCAGGAGACATCCAACCTATGAATACTCATTTGAGCGTCTCTT
GTCTAAGTTGACAAGCAATGTTCAA

SP070 amino acid (SEQ ID NO:116)

HQMGHKVQGSDEKYYFTQRGLEQAGITILPFDEKNLDGDMETIAGNAFRPDNNVEIAYADQNGISYKR
YHEFLGSFMRDFVSMGVAGAHGKTSTTGMLSHVLSHITDTSFLIGDGTGRGSANAKYFVFESDEYERHF
MPYHPEYSIITNIDFDHPDYFTSLEDVFNAFNDYAKQITKGLFVYGEDAELRKITSDAPIYYYGFEEAG
NDFVASDLLRSITGSTFTVHFRGNLQGFHIPTFGRHNIMNATAVIGLLYTAGFDLNLVREHLKTFAGV
KRRFTEKIVNDTVIIDDFAHHPTEIIATLDAARQKYPKSKEIVAVFQPHFTFTRTIALLLDDFAHALNQADA
VYLAQIYGSAREVDHGDVKVEDLANKINKKHQVITVENVSPLLDHNAVYVFMGAGDIQTYEYSFERLL
SNLTSNVQ

SP071 nucleotide (SEQ ID NO:117)

TTTTAACCCAACTGTTGGTACTTTCTTTTTACTGCAGGATTGAGCTTGTTAGTTTTATTGGTTTTCTAA
AAGGGAAAATGGAAGAAACGACTTGTTTCATTTTCTGCTGTTGACTAGCATGGGAGTTCAATTGTTGCC
GGCCAGTGCTTTTGGGTTGACCAGCCAGATTTTATCTGCCTATAATAGTCAGCTTTCTATCGGAGTCGG
GGAACATTTACCAGAGCCTCTGAAAATCGAAGGTTATCAATATATTGGTTATATCAAAACTAAGAAACA
GGATAATACAGAGCTTTCAAGGACAGTTGATGGGAAATACTCTGCTCAAAGAGATAGTCAACCAAACCTC
TACAAAACATCAGATGTAGTTCATTCAGCTGATTTAGAATGGAACCAAGGACAGGGGAAGGTTAGTTT
ACAAGGTGAAGCATCAGGGGATGATGGACTTTCAGAAAAATCTTCTATAGCAGCAGACAATCTATCTTC
TAATGATTTCATTCGCAAGTCAAGTTGAGCAGAATCCGGATCACAAAGGAGAATCTGTAGTTCGACCAAC
AGTGCCAGAACAAGGAAATCCTGTGTCTGTCTACAAACGGTGCAGAGTGCAGGAAGAGGAAGTATTGGCGAC
GACAAATGATCGACCAGAGTATAAACTTCCATTGGAAACCAAGGCACGCAAGAACCCGGTTCATGAGGG
TGAAGCCGCGAGTCCGTGAAGACTTACCAGTCTACACTAAGCCACTAGAAACCAAGGTACACAAGGACC
CGGACATGAAGGTGAAGCTGCAGTTCGCGAGGAAGAACCAGCTTACACAGAACCGTTAGCAACGAAAGG
CACGCAAGAGCCAGGTCATGAGGGCAAAGCTACAGTCCGCGAAGAGACTCTAGAGTACACGGAACCGGT
AGCGACAAAAGGCACACAAGAACCCGAACATGAGGGCGAaCGGsCAGTAGAAGAAGAACTTCCGGCTTT
AGAGGTCACTACACGAAATAGAACGGAATCCAGAATATTCCTTATACAACAGAAGAAATTCAGGATCC
AACACTTCTGAAAAATCGTCGTAAGATTGAACGACAAGGGCAAGCAGGGACACGTACAATTCATATGA
AGACTACATCGTAAATGGTAATGTCTGTAGAACTAAAGAAGTGTACGAACCTGAAGTAGCTCCGGTCAA
CGAAGTCGTTAAAGTAGGAACACTTGTGAAAAGTTAAACCTACAGTAGAAATTACAAACTTAAACAAAAGT
TGAGAACAAAAAATCTATAACTGTAAGTTATAACTTAATAGACACTACCTCAGCATATGTTTCTGCAAA
AACGCAAGTTTTCCATGGAGACAAGCTAGTTAAAGAGGTGGATATAGAAAATCCTGCCAAAGAGCAAGT
AATATCAGGTTTAGATTACTACACACCGTATACAGTTAAAACACACCTAACTTATAATTTGGGTGAAAA
TAATGAGGAAAATACTGAAACATCAACTCAAGATTTCCAATTAGAGTATAAGAAAATAGAGATTAAAGA
TATTGATTTCAGTAAATTATACGGTAAAGAAAATGATCGTTATCGTAGATATTTAAGTCTAAGTGAAGC
GCCGACTGATACGGCTAAATACTTTGTAAAAGTGAAATCAGATCGCTTCAAAGAAATGTACCTACCTGT
AAAATCTATTACAGAAAATACGGATGGAACGTATAAAGTGACGGTAGCCGTTGATCAACTTGTTCGAAGA
AGGTACAGACGGTTACAAAGATGATTACACATTTACTGTAGCTAAATCTAAAGCAGAGCAACCAGGAGT
TTACACATCCTTTAAACAGCTGGTAACAGCCATGCAAAGCAATCTGTCTGGTGTCTATACATTGGCTTC
AGATATGACCGCAGATGAGGTGAGCTTAGGCGATAAGCAGACAAGTTATCTCAGAGGTGCATTTACAGG
GAGCTTGATCGGTTCTGATGGAACAAAATCGTATGCCATTTATGATTTGAAGAAACCATTATTTGATAC
ATTAAATGGTGCTACAGTTAGAGATTTGGATATTAATACTGTTTCTGCTGATAGTAAAGAAAATGTCGC
AGCGCTGGCGAAGGCAGCGAATAGCGCGAATATTAATAATGTTGCAGTAGAAGGAAAAATCTCAGGTGC
GAAATCTGTTGCGGGATTAGTAGCGAGCGCAACAAAATACAGTGATAGAAAACAGCTCGTTTACAGGGAA
ACTTATCGCAAATCACCAGGACAGTAATAAAAATGATACTGGAGGAATAGTAGGTAATATAACAGGAAA
TAGTTTCGAGAGTTAATAAAGTTAGGGTAGATGCCCTTAATCTCTACTAATGCACGCAATAATAACCAAAC
AGCTGGAGGGGATAGTAGGTAGATTAGAAAATGGTGCATTGATATCTAATTCGGTTGCTACTGGAGAAAAT
ACGAAATGGTCAAGGATATTCTAGAGTCGGAGGAATAGTAGGATCTACGTGGCAAAACGGTCGAGTAAA
TAATGTTGTGAGTAACGTAGATGTTGGAGATGGTTATGTTATCACCGGTGATCAATACGCAGCAGCAGA
TGTGAAAAATGCAAGTACATCAGTTGATAATAGAAAAGCAGACAGATTGCTACAAAATTATCAAAAAGA
CCAAATAGACGCGAAAGTTGCTGATTATGGAATCACAGTAACTCTTGATGATACTGGGCAAGATTTAAA
ACGTAATCTAAGAGAAGTTGATTATACAAGACTAAATAAAGCAGAAGCTGAAAGAAAAGTAGCTTATAG
CAACATAGAAAAACTGATGCCATTCTACAATAAAGACCTAGTAGTTCACTATGGTAACAAAGTAGCGAC
AACAGATAAACTTTACACTACAGAATTGTTAGATGTTGTGCCGATGAAAGATGATGAAGTAGTAACGGA
TATTAATAATAAGAAAAATTCAATAAATAAAGTTATGTTTACATTTCAAAGATAATACAGTAGAATACCT
AGATGTAACATTCAAAGAAAACCTTCATAAACAGTCAAGTAATCGAATACAATGTTACAGGAAAAAGATA
TATATTCACACCAGAAGCATTGTTTCAGACTATACAGCGATAACGAATAACGTACTAAGCGACTTGCA
AAATGTAACACTTAAC

SP071 amino acid (SEQ ID NO:118)

FNPTVGTFLFTAGLSLLVLLVSKRENGKKRLVHFLLLTSMGVQLLPASAFGLTSQILSAYNSQLSIGVG
EHLPEPLKIEGYQYIGYIKTKKQDNTELSRTVDGKYSAQRDSQPNSTKTSDVVHSADLEWNQGQGVSL
QGEASGDDGLSEKSSIAADNLSSNDSFASQVEQNPDHKGESVVRPTVPEQGNPVSATTVQSAEEEEVLAT
TNDRPEYKLPLETKGTQEPGHEGEAAVREDLPVYTKPLETKGTQGPGEHEGEAAVREEEPAYTEPLATKG
TQEPGHEGKATVREETLEYTEPVATKGTQEPEHEGERXVEEELPALEVTTRNRTEIQNI PYTTEEIQDP
TLLKNRRKIERQGQAGTRTIQYEDYIVNGNVVETKEVSRTEVAPVNEVVKVGTLVKVKPTVEITNLTKV
ENKKSITVSYNLIDTTSAYVSAKTQVFHGDKLVEVDIENPAKEQVISGLDYTPYTVKTHLTYNLGEN
NEENTETSTQDFQLEYKKIEIKDIDSVELYGKENDRYRRYLSLSEAPTDTAKYFVKVKSDFRKEMYLPV
KSITENTDGTYKVTVAVDQLVEEGTDGYKDDYTFTVAKSKAEQPGVYTSFKQLVTAMQSNLSGVYTLAS
DMTAEVSLGDKQTSYLTGAFTGSLIGSDGTSYAIYDLKKPLFDTLNGATVRDLDIKTVSADSKENVA
ALAKAANSANINNVAVEGKISGAKSVAGLVASATNTVIENSSFTGKLIANHQDSNKNDTGGIVGNITGN
SSRVNKVRVDALISTNARNNNQTAGGIVGRLENGALISNSVATGEIRNGQGYSRVGGIVGSTWQNGRVN
NVVSNVDVG DG YVITGDQYAAADVKNASTVDNRKADRFATKLSKDQIDAKVADYGITVTLDDTGQDLK
RNLREVDYTRLNKAEAEKRVAYSNIEKLMPFYNKDLVVHYGNKVATTDKLYTTELLDVVPMKDDEVTD
INNKKNSINKVMLHFKDNTVEYLDVTFKENFINSQVIEYNVTGKEYIFTPEAFVSDYTAITNNVLSDLQ
NNNLTN

SP072 nucleotide (SEQ ID NO:119)

TTTTTAACCCAACACTGTTGGTACTTTTCCTTTTACTGTCAGGATTAGACTTGTAGTTTATTGGTTTCTAA
AAGGGAAAATGGAAGAAACGACTTGTTTCATTTTCTGCTGTTGACTAGCATGGGAGTTCAATTGTTGCC
GGCCAGTGCTTTTGGGTTGACCAGCCAGATTTTATCTGCCTATAATAGTCAGCTTCTATCGGAGTCGG
GGAACATTTACCAGAGCCTCTGAAAATCGAAGGTTATCAATATATTGGTTATATCAAAACCTAAGAAACA
GGATAATACAGAGCTTTCAAGGACAGTTGATGGGAAATACTCTGCTCAAAGAGATAGTCAACCAAACTC
TACAAAACATCAGATGTAGTTCATTTCAGCTGATTTAGAATGGAACCAAGGACAGGGGAAGGTTAGTTT
ACAAGGTGAAGCATCAGGGGATGATGGACTTTCAGAAAAATCTTCTATAGCAGCAGACAATCTATCTTC
TAATGATTTCATTTCGCAAGTCAAGTTGAGCAGAATCCGGATCACAAGGAGAATCTGTAGTTCGACCAAC
AGTGCCAGAACAAGGAAATCCTGTGTCTGCTACAACGGTGCAGAGTGCGGAAGAGGAAGTATTGGCGAC
GACAAATGATCGACCAGAGTATAAACTTCCATTGGAAACCAAAGGCACGCAAGAACCCGGTCATGAGGG
TGAAGCCGCAGTCCGTGAAGACTTACCAGTCTACACTAAGCCACTAGAAACCAAAGGTACACAAGGACC
CGGACATGAAGGTGAAGCTGCAGTTTCGCGAGGAAGAACCAGCTTACACAGAACCCTTAGCAACGAAAGG
CACGCAAGAGCCAGGTTCATGAGGGCAAAGCTACAGTCCGCGAAGAGACTCTAGAGTACACGGAACCGGT
AGCGACAAAAGGCACACAAGAACCCGAACATGAGGGCGAaCGGsCAGTAGAAGAAGAACTTCCGGCTTT
AGAGGTCACTACACGAAATAGAACGGAAATCCAGAATATTCCTTATACAACAGAAGAAATTCAGGATCC
AACACTTCTGAAAAATCGTCGTAAGATTGAACGACAAGGGCAAGCAGGGACACGTACAATTCAATATGA
AGACTACATCGTAAATGGTAATGTCTGTAGAACTAAAGAAGTGTCACGAACTGAAGTAGCTCCGGTCAA
CGAAGTCGTTAAAGTAGGAACACTTGTGAAAGTTAAACCTACAGTAGAAATTACAACTTAACAAAAGT
TGAGAACAAAAATCTATAACTGTAAGTTATAACTTAATAGACACTACCTCAGCATATGTTTCTGCAAA
AACGCAAGTTTTTCATGGAGACAAAGCTAGTTAAAGAGGTGGATATAGAAAATCCTGCCAAAGAGCAAGT
AATATCAGGTTTTAGATTACTATACACACCTGATATACAGTTAAACACACCTAACTTATAATTTGGGTGAAAA
TAATGAGGAAAAATCTGAAACATCAACTCAAGATTTCCAATTAGAGTATAAGAAAAATAGAGATTAAAGA
TATTGATTTCAGTAGAATTATACGGTAAAGAAAAATGATCGTTATCTGTAGA

SP072 amino acid (SEO ID NO:120)

FNPTVGTFLFTAGLSLLVLLVSKRENGKKRLVHFLLLTSMGVQLLPASAFGLTSQILSAYNSQLSIGVG
EHLPEPLKIEGYQYIGYIKTKKQDNTELSRTVDGKYSAQRDSQPNSTKTSADVHSADLEWNQGGKVS
QGEASGDDGLSEKSSIAADNLSSNDSFASQVEQNPDPHGKESVVRPTVPEQGNPVSATTVQSAAEEVLAT
TNDRPEYKLPLETGKTQEPGHEGEAAVREDLPVYTKPLETGTGQPGHEGEAAVREEEPAYTEPLATKG
TQEPGHEGKATVREETLEYTEPVATKGTQEPHEGERXVEELPALEVTTRNRTEIQNIPYTTEEIQDP
TLLKNRRKIERQGQAGTRTIQYEDYIVNGNVVETKEVSRTVAPVNEVVKVGTLVKVKPTVEITNLTKV
ENKKSITVSYNLIDTTSAYVSAKTQVFHGDGLVKEVDIENPAKEQVISGLDYYTPYTVKTHLTYNLGEN
NEENTETSTODFOLEYKKIEIKDIDSVELYGKENDRYRR

SP073 nucleotide (SEQ ID NO:121)

TCGTAGATATT'TAAGTCTAAGTGAAGCGCCGACTGATACGGCTAAATACCTTTGTAAAAGTGAAATCAGATCGCTTCAAAGAAATGTACCTACCTGTAAAATCTATTACAGAAAATACGGATGGAACGTATAAAGTGACGGTAGCCGTTGTATCAACCTTGTGCGAAGAAAGGTACAGACGGTTACAAAGATGATTACACATTTACTGTAGCTAAATCTAAAGCAGACGACCAACCGAGGTTTACACATCTTTTAAACAGCTGGTATCAACGCCATGCAACGACAACTCTGCTGGTGCTATACATATTGGCTTCAGATATGACCGCAGATGAGGTGAGCTTAGGCCGATAAGCAGAC

SECRET

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AAGTTATCTCACAGGTGCATTTACAGGGAGCTTGATCGGTTCTGATGGAACAAAATCGTATGCCATTTA
TGATTTGAAGAAACCATTTATTTGATACATTAAATGGTGCTACAGTTAGAGATTTGGATATTAAACCTGT
TTCTGCTGATAGTAAAGAAAATGTGCGAGCGCTGGCGAAGGCAGCGAATAGCGCGAATATTAATAATGT
TGCAGTAGAAGGAAAAATCTCAGGTGCGAAATCTGTTGCGGGATTAGTAGCGAGCGCAACAAAATACAGT
GATAGAAAACAGCTCGTTTACAGGGAAACTTATCGCAAATCACCAGGACAGTAATAAAAAATGATACTGG
AGGAATAGTAGGTAATATAACAGGAAATAGTTCGAGAGTTAATAAAGTTAGGGTAGATGCCCTTAATCTC
TACTAATGCACGCAATAATAACCAAACAGCTGGAGGGATAGTAGGTAGATTAGAAAATGGTGCATTGAT
ATCTAATTCGGTTGCTACTGGAGAAAATACGAAATGGTCAAGGATATTCTAGAGTCGGAGGAATAGTAGG
ATCTACGTGGCAAACGGTCGAGTAAATAATGTTGTGAGTAACGTAGATGTTGGAGATGGTTATGTTAT
CACCGGTGATCAATACGCAGCAGCAGATGTGAAAAATGCAAGTACATCAGTTGATAATAGAAAAGCAGA
CAGATTCGCTACAAAATTATCAAAAGACCAAATAGACGCGAAAGTTGCTGATTATGGAATCACAGTAAC
TCTTGATGATACTGGGCAAGATTTAAACGTAATCTAAGAGAAGTTGATTATACAAGACTAAAATAAAGC
AGAAGCTGAAAGAAAAGTAGCTTATAGCAACATAGAAAACTGATGCCATTCTACAATAAAGACCTAGT
AGTTCACATATGGTAACAAAGTAGCGACAACAGATAAACTTTACACTACAGAATTGTTAGATGTTGTGCC
GATGAAAGATGATGAAGTAGTAACGGATATTAATAATAAGAAAAATTCATAAAATAAAGTTATGTTACA
TTTCAAAGATAATACAGTAGAATACCTAGATGTAACATTCAAAGAAAACCTTCATAAACAGTCAAGTAAT
CGAATACAATGTTACAGGAAAAGAATATATATTCACACCAGAAGCATTGTTTCAGACTATACAGCGAT
AACGAATAACGTACTAAGCGACTTGCAAAATGTAACACTTAAC

SP073 amino acid (SEQ ID NO:122)

RRYLSLSEAPTDKAKYFVKVKSDFKEMYPVKISITENTDGTIKVTVAVDQLVVEEGTDGYKDDYTFTVA
KSKAEQPGVYTSFKQLVTAMQSNLSGVYTLASDMTAEVSLGDKQTSYLTGAFTGSLIGSDGTSYAIY
DLKKPLFDTLNGATVRDLDIKTVSADSKENVAALAKAANSANINNVAVEGKISGAKSVAGLVASATNTV
IENSSTGKLIANHQDSNKNNDTGGIVGNITGNSSRVNKVRVDALISTNARNNNQTAGGIVGRLENGALI
SNSVATGEIRNGQYSRVGGIVGSTWQNGRVNNVSNVDVGDGYVITGDQYAAADVKNASTVDNRKAD
RFATKLSKDQIDAKVADYGITVTLDDTGQDLKRNLEVDYTRLNKAERKVAYSNIEKLMPFYNKDLV
VHYGNKVATTDKLYTTELLDVVPMKDDEVVTDINNKKNSINKVMLHFKDNTVEYLDVTFKENFINSQVI
EYNVTGKEYIFTPEAFVSDYTAITNNVLSDLQNVTLN

SP074 nucleotide (SEQ ID NO:123)

CTTTGGTTTTGAAGGAAGTAAGCGTGACAATTTGCTGTAGAAAGGAATCAATCAACTTCGTGAGCATGT
AGACACTCTATTGATTATCTCAAACAACAATTTGCTTGAAATTGTTGATAAGAAAACACCGCTTTTGGA
GGCTCTTAGCGAAGCGGATAACGTTCTTCGTCAAGGTGTTCAAGGGATTACCGATTTGATTACCAATCC
AGGATTGATTAACTTTGACTTTGCCGATGTGAAAACGGTAATGGCAAACAAAGGAATGCTCTTATGGG
TATTGGTATCGGTAGTGGAGAAGAACGTGTGGTAGAAGCGGCACGTAAGGCAATCTATTACCACCTTCT
TGAAACAACATATTGACGGTGCTGAGGATGTTATCGTCAACGTTACTGGTGGTCTTGACTTAACCTTGAT
TGAGGCAGAAGAGGCTTCACAAATTGTGAACCAGGCAGCAGGTCAAGGAGTGAACATCTGGCTCGGTAC
TTCAATTGATGAAAGTATGCGTGATGAAATTCGTGTAACAGTTGTTGCAACGGGTGTTCTGTCAGACCG
CGTAGAAAAGGTTGTGGCTCCACAAGCTAGATCTGCTACTAACCCTGAGACAGTGAAACCAGCTCA
TTCACATGGCTTTGATCGTCATTTTGATATGGCAGAAACAGTTGAATTGCCAAAACAAAATCCACGTCG
TTTGGAACCAACTCAGGCATCTGCTTTTGTTGATTGGGATCTTCGCCGTGAATCGATTGTTCTGTACAAC
AGATTCACTCGTTTCTCCAGTCGAGCGCTTTGAAGCCCCAATTTACAAGATGAAGATGAATTGGATAC
ACCTCCATTTTTTCAAAAATCGT

SP074 amino acid (SEQ ID NO:124)

FGFEGSKRGQFAVEGINQLREHVDTLIIISNNLLEIVDKKTPLEALSEADNVLRQGVQGITDLITNP
GLINLDFADVKTVMANKGNALMGIGIGSGEERVVEAARKAIYSPLLETTIDGAEDVIVNVTGGGLDLTLI
EAEESQIVNQAAAGQGVNIWLGTSIDESMRDEIRVTVVATGVRQDRVEKVVAPQARSATNYRETVKPAH
SHGFDRHFDMAETVELPKQNPRLLEPTQASAFGDWDLRRESIVRTTDSVSVSPVERFEAPISQDEDELDT
PPFFKNR

SP075 nucleotide (SEQ ID NO:125)

CTACTACCTCTCGAGAGAAAGTGACCTAGAGGTGACCGTTTTTGACCATGAGCAAGGTCAAGCCACCAA
GGCCGCAGCAGGAATTATCAGTCCTTGGTTTTCCAAACGCCGTAATAAAGCCTGGTACAAGATGGCGCG
CTTGGGGGCTGATTTTTATGTGGATTTATTAGCTGATTTAGAGAAATCAGGACAAGAAATCGACTTTTA
CCAGCGTTCGGGAGTCTTTCTCTTGAAAAAGGATGAATCCAATTTGGAAGAAGTTTATCAACTGGCCCT
CCAGCGCAGAGAAGAATCTCCCTTGATAGGGCAATTAGCCATTCTGAACCAAGCCTCAGCTAATGAATT
ATTCCCTGGTTTTGCAGGGATTTGACCGCCTGCTCTATGCTTCTGGTGGAGCGAGAGTAGATGGCCAACT

TTTAGTGA CTGCTTTGCTGGAAGTCAGTCATGTCAAGCTGGTCAAAGAAAAAGTGACTCTGACACCGTT
AGCATCAGGCTACCAGATTGGTGAAGAGGAGTTTGAGCAGGTTATTTTGGCGACGGGAGCTTGGTTGGG
GGACATGTTAGAGCCTTTAGGTTATGAAGTGGATGTCCGTCTCAAAAAGGACAACCTACGAGATTATCA
GCTTGCCCCAAGACATGGAAGATTACCCTGTTGTCATGCCAGAAGGGGAGTGGGATTTGATTCCCTTTGC
AGGTGGGAAATTATCCTTAGGCGCTACCCACGAAAATGACATGGGATTTGATTTGACGGTAGATGAAAC
CTTGCTCCAACAAATGGAGGAGGCCACCTTGACTCACTATCTGATTTTGGCTGAAGCTACTTCAAATC
TGAGCGTGTGGAATCCGTGCCTACACCAGTGATTTCTCTCCTTTCTTTGGGCAGGTGCCTGACTTAAC
TGGTGTCTATGCAGCCAGTGGACTAGGTTTCATCAGGCCTCACAACCTGGTCTATCATTTGGTTACCATCT
AGCCCAACTGATCCAAGACAAGGAGTTGACCTTGGACCCTCTAAATTACCCAATTGAAAACCTATGTCAA
ACGAGTAAAAAGCGAA

SP075 amino acid (SEQ ID NO:126)

YYLSRES DLEVTVFDHEQ GQATKAAAGIISPWF SKRRNKAWYKMARLGADFYVDLLADLEKSGQEIDFY
QRSGVFL LKKDESNLEELYQLALQRREESPLIGQLAILNQASANELFPGLQGFDRLLYASGGARVDGQL
LVTRLLEVSHVKLVKEKVTLTPLASGYQIGEEEFQVILATGAWLGDMLEPLGYEVDVRPQKGQLRDYQ
LAQDMEDY PVMPEGEWDLIPFAGGKLSLGATHENDMGFDLTVDETLLQQMEEATLTHYLILAEATSKS
ERVGIRAYTSDFS PFFGQV PDLTGVYAASGLSSGLTTGPIIGYHLAQLIQDKELTLDPLNYPINENYVK
RVKSE

SP076 nucleotide (SEQ ID NO:127)

TAAGGTCAA AAGTCAGACCGCTAAGAAAGTGCTAGAAAAGATTGGAGCTGACTCGGTTATCTCGCCAGA
GTATGAAATGGGGCAGTCTCTAGCACAGACCATTCTTTTCCATAATAGTGTTGATGTCTTTCAGTTGGA
TAAAAATGTGTCTATCGTGGAGATGAAAATTCCTCAGTCTTGGGCAGGTCAAAGTCTGAGTAAATTAGA
CCTCCGTGGCAAATACAATCTGAATATTTTGGGTTTCCGAGAGCAGGAAAATTCCTCATTTGGATGTTGA
ATTTGGACCAGATGACCTCTTGAAAGCAGATACCTATATTTTGGCAGTCATCAACAACCAGTATTTGGA
TACCCTA

SP076 amino acid (SEQ ID NO:128)

KVKSQTA KKVLEKIGADSVISPEYEMGQSLAQ TILFHNSVDVFQ LDKNVSIVEMKIPQSWAGQSLSKLD
LRGKYNL NILGFREQENSPLDVEFGPDDL LKADTYILAVINNQYLDTL

SP077 nucleotide (SEQ ID NO:129)

TGACGGGTCTCAGGATCAGACTCAGGAAATCGCTGAGTGTTAGCTAGCAAGTATCCTAATATCGTTAG
AGCCATCTATCAGGAAAATAAATGCCATGGCGGTGCGGTCAATCGTGGCTTGGTAGAGGCTTCTGGGCG
CTATTTTAAAGTAGTTGACAGTGACTGGGTGGATCCTCGTGCCTACTTGAAAATTCCTGAAACTTG
CAGGAACCTTGAGAGCAAAGGTCAAGAGGTGGATGTCTTTG

SP077 amino acid (SEQ ID NO:130)

DGSQDQTQ EIAECLASKYPNIVRAIYQENKCHGGAVNRGLVEASGRYFKVVDSDDWVDP RAYLKILETC
RNLRAKVKRWMSL

SP078 nucleotide (SEQ ID NO:131)

TAGAGGCTTTGCCAAATGGTGGGAAGGGCACGAGCGTCGAAAAGAGGAACGCTTTGTCAAACAAGAAGA
AAAAGCTCGCCAAAAGGCTGAGAAAAGAGGCTAGATTAGAACAAGAAGAGACTGAAAAAGCCTTACTCGA
TTTGCCCTCCTGTTGATATGGAACGGGTGAAATTCTGACAGAGGAAGCTGTTCAAATCTTCCACCTAT
TCCAGAAGAAAAGTGGGTGGAACAGAAAATCATCCTGCCTCAAGCTGAACTTAAATTCCTGAAACAGGA
AGATGACTCAGATGACGAAGATGTTTCAGGTCGATTTTTCAGCCAAAGAAGCCCTTGAATACAACTTCC
AAGCTTACAACCTTTTGCACCAGATAAAACCAAAAGATCAGTCTAAAGAGAAGAAAATTGTCAGAGAAAA
TATCAAAATCTTAGAAGCAACCTTTGCTAGCTTTGGTATTAAGGTAACAGTTGAACGGGCGGAAATTGG
GCCATCAGTGACCAAGTATGAAGTCAAGCCGGCTGTTGGTGTAAAGGTCAACCGCATTTCCAATCTATC
AGATGACCTCGCTAGCCTTGGCTGCCAAAGATGTCGGATTGAAGCACC AATCCCTGGGAAATCCCT
AATCGGAATTGAAGTGCCCAACTCCGATATTGCCACTGTATCTTTCCGAGA ACTATGGGAACAATCGCA
AACGAAAGCAGAAAATTTCTTGAAAATTCCTTAGGGAAGGCTGTTAATGGAACCGCAAGAGCTTTTGA
CCTTTCTAAAATGCCCCACTTGCTAGTTGCAGGTTCAACGGGTTCAAGGAAGTCAGTAGCAGTTAACGG
CATTATTGCTAGCATTCTCATGAAGGCGAGACCAGATCAAGTTAAATTTATGATGGTTCGATCCCAAGAT
GGTTGAGTTATCTGTTTACAATGATATTCCCCACCTCTTGATTCCAGTCGTGACCAATCCACGCAAAGC
CAGCAAGGCTCTGCAAAGGTTGTGGATGAAATGGAACCGTTATGAACTCTTTGCCAAGGTGGGAGT
TCGGAATATTGCAGGTTTTAATGCCAAGGTAGAAGAGTTCAATTCCAGTCTGAGTACAAGCAAATTC

GCTACCATTTCATTGTCGTGATTGTGGATGAGTTGGCTGACCTCATGATGGTGGCCAGCAAGGAAGTGGAGATGCTATCATCCGCTCTGGGCAGAAGGCGCGTGCAGGTATCCACATGATTCTTGCAACTCAGCGTCCATCTGTTGATGTCATCTCTGGTTTGATTAAAGGCCAATGTTCCATCTCGTGTAGCATTTCGCGTTTCATCAGGAACAGACTCCCGTACGATTTTGGATGAAAATGGAGCAGAAAACTTCTTGGTCGAGGAGACATGCTCTTTAAACCGATTGATGAAAATCATCCAGTTCGCTCTCCAAGGCTCCTTTATCTCGGATGACGATGTGAGCGCATTGTGAACCTTCATCAAGACTCAGGCAGATGCAGACTACGATGAGAGTTTTGATCCAGGTGAGGTTTCTGAAAATGAAGGAGAATTTTCGGATGGAGATGCTGGTGGTGTATCCGCTTTTTGAAGAAGCTAAGTCTTTGGTTATCGAAACACAGAAAGCCAGTGCCTCTATGATTACGCGTCGTTTATCAGTTGGATTTAACCGTGCACCCGCTCTCATGGAAGAACTGGAGATAGCAGGTGTCATCGGTCCAGCTGAAGGTACCAACC

SP078 amino acid (SEQ ID NO:132)

RGFAKWEGHERRKEERFVKQEEKARQKAEKEARLEQEETEKALLDLPPVDMETGEIILTEEAVQNLPPPIPEEKWVEPEIILPQAEKLFPEQEDSDDEDVQVDFSAKEALEYKLPQLFAPDKPKDQSKEKKIVREN IKILEATFASFGIKVTVRAEIGPSVTKYEVKPAVGVRVNRISNLSDDLALALAAKDVRIEAPIPGKSLIGIEVPNSDIATVSFRELWEQSQTAEENFLEIPLGKAVNGTARAFDL SKMPHLLVAGSTGSGKSVAVNG I IASILMKARPDQVKFMMVDPKMVELSVYNDI PHLLIPVVTNPRKASKALQKVVDENRYELFAKVGVRNIAGFNAKVEEFNSQSEYKQIPLPFIIVVIVDELADLMMVASKEVEDAIIRLGQKARAAGIHMILATQRP SVDVISGLIKANVPSRVAFVSSGTDSTILDENGAEKLLGRGDMFLFKPIDENHPVRLQGSFISDDDV ERIVNFIKTQADADYDESFDPGEVSENEGEFSDGDAGDPLFEEAKSLVIETQKASASMIQRRLSVGFNRATRLMEELEIAGVIGPAEGTKPRKVLQQ

SP079 nucleotide (SEQ ID NO:133)

TCAAAAAGAGAAGGAAAACCTTGGTTATTGCTGGGAAAATAGGTCCAGAACCAGAAAATTTGGCCAATATGTATAAGTTGCTGATTGAAGAAAATACCAGCATGACTGCGACTGTTAAACCGAATTTTGGGAAGACAAGCTTCCTTTTATGAAGCTCTGAAAAAAGGCGATATTGACATCTATCCTGAATTTACTGGTACGGTGACTGA AAGTTTGCTTCAACCATCACCAAGGTGAGTCATGAACCAGAACAGGTTTATCAGGTGGCGCGTGATGGCATTGCTAAGCAGGATCATCTAGCCTATCTCAAACCCATGTCTTATCAAAACACCTATGCTGTAGCTGT TCCGAAAAAGATTGCTCAAGAATATGGCTTGAAGACCATTTCAGACTTGAAAAAGTGGAAAGGCGAGTTGAAGGCAGGTTTTACACTCGAGTTTAACGACCCGTGAAGATGGAAATAAGGGCTTGCAATCAATGTATGG TCTCAATCTCAATGTAGCGACCATTGAGCCAGCCCTTCGCTATCAGGCTATTACAGTCAGGGGATATTCA AATCACGGATGCCTATTTCGACTGATGCGGAATTGGAGCGTTATGATTTACAGGTCTTGGAAGATGACAA GCAACTCTTCCCACCTTATCAAGGGGCTCCACTCATGAAAGAAGCTCTTCTCAAGAAACACCCAGAGTT GGAAAGAGTTCTTAATACATTGGCTGGTAAGATTACAGAAAGCCAGATGAGCCAGCTCAACTACCAAGT CGGTGTTGAAGGCAAGTCAGCAAAGCAAGTAGCCAAGGAGTTTCTCCAAGAACAAAGGTTTGTGTAAGAA A

SP079 amino acid (SEQ ID NO:134)

QKEKENLVIAGKIGPEPEILANMYKLLIEENTSMATVKNPFGKTSFLYEALKKGDIDIDYPEFTGTVTE SLLQPSPKVSHEPEQVYQVARDGIAKQDHLAYLKPMSYQNTYAVAVPKKIAQEQYGLKTI SDLKKVEGQL KAGFTLEFNDREDGNKGLQSMYGLNLNVATIEPALRYQAIQSGDIQITDAYSTDAELERYDLQVLEDDK QLFPPYQGAAPLMKEALLKKHPELERNLNTLAGKITESQMSQLNYQVGVEGKSAKQVAKEFLQEQGLLKK

SP080 nucleotide (SEQ ID NO:135)

ACGTTCTATTGAGGACCACTTTGATTCAAACCTTCGAATTGGAATATAACCTCAAAGAAAAAGGGAAAACAGATCTTTTGAAGCTAGTTGATAAAACAACGACATGCGTCTGCATTTTATCCGCCAAACTCATCCACG CGGTCTCGGAGATGCTGTTTTGCAAGCCAAGGCTTTCGTCGGAAATGAACCTTTTGTCTGTTATGCTTGG TGATGACTTGATGGATATCACAGACGAAAAGGCTGTTCCACTTACCAAACAACCTCATGGATGACTACGAGCGTACCCACGCGTCTACTATCGCTGTATGCCAGTCCCTCATGACGAAGTATCTGCTTACGGGGTTAT TGCTCCGCAAGGCGAAGGAAAAGATGGTCTTTACAGTGTGTTGAAACCTTTGTTGAAAAACCAGCTCCAGAGGACGCTCCTAGCGACCTTGCTATTATCGGACGCTACCTCCACGCTGAAATTTTGAGATTCTCGA AAAGCAAGCTCCAGGTGCAGGAAATGAAATTCAGCTGACAGATGCAATCGACACCCTCAATAAAACACA ACGTGTATTTGCTCGTGAGTTCAAAGGGGCTCGTTACGATGTCGGAGACAAGTTTGGCTTCATGAAAAC ATCCATCGACTACGCCCTCAAACACCCACAAGTCAAAGATGATTTGAAGAATTACCTCATCCAACCTTGG AAAAGAATTGACTGAGAAGGAA

SP080 amino acid (SEQ ID NO:136)

RSIEDHFDNSFELEYNLKEKGKTDLLKLVDKTTDMRLHFIRQTHPRGLGDAVLQAKAFVGNPEFVVMLG
DDLMDITDEKAVPLTKQLMDDYERTHASTIAVMPVPHDEVSAYGVIAPQGEGKDGLYSVETFVEKPAP
DAPSDLAIIGRYLLTPEIFEILEKQAPGAGNEIQLTDAIDTLNKTQRVFAREFKGARYDVGDKFGFMKT
SIDYALKHPQVKDDLKNYLIQLGKELTEKE

SP081 nucleotide (SEQ ID NO:137)

CGCTCAAATACCAGAGGTGTTTCAGCTAATCGAGCACGTTTCTCCTCAAATGTTGAAAGCCCAATTGGA
GAGTGTCTTTTCTGATATTCCACCTCAGGCTGTAAAACTGGAATGTTGGCTACTACTGAAATCATGGA
AATCATCCAACCCTATCTTAAAAAAGTGGATTGTCCCTATGTCCTTGATCCTGTTATGGTTGCTACAAG
TGGAGATGCCTTGATTGACTCAAATGCTAGAGACTATCTCAAAACAACTTACTACCTCTAGCAACTAT
TATTACGCCAAATCTTCTGAAGCAGAAAGAGATTGTTGGTTTTCAATCCATGACCCCGAAGACATGCA
GCGTGTGGTGCCTGATTTTAAAGAATTTGGTCTCAGTCTGTGGTTATCAAGGCGGACATCTCAA
AGGTGGTGTAAAGATTTCCTCTTTACCAAGAATGAACAATTTGTCTGGGAAAGCCCACGAATTCAAAC
CTGTCACACCCATGGTACT

SP081 amino acid (SEQ ID NO:138)

AQNTRGVQLIEHVSPQMLKAQLESVFS DIPQAVKTGMLATTEIMEIIQPYLKKLDPCPYVLDPVMVATS
GDALIDS NARDYLKTNLLPLATIIITPNLPEAEIIVGFSIHDPEDMQRAGRLLILKEFGPQSVVIKGGHLK
GGAKDFLFTKNEQFVWESPRIQTCHTHGT

SP082 nucleotide (SEQ ID NO:139)

AATTGTACAATTAGAAAAAGATAGCAAATCAGACAAAGAACAAGTTGATAAACTATTTGAATCATTGGA
TGCATCTTCAGATGAATCTATTTCTAAATTAAGAAGTATCTGAACTTCACTTAAACCGATGCAGG
TAAAGACTATCTTAATAACAAAGTCAAAGAATCATCTAAAGCAATTGTAGATTTTCATTGCAAAAAGG
TTTGGCTTATGATGTTAAAGATTGAGATGACAAATTTAAAGATAAAGCAACTCTTGAAACAAATGTAAA
AGAAATTACAAACAAATTGATTTTATCAAAAAAGTTGATGAACTTTTAAACAAGAGAATTTGGAAGA
AACTCTTAAATCTCTAAATGATCTTGTGTTGATAAATATCAAAACAAATCGAACTTTTGAAGAAAAGA
AGAAAAAGCTGCTGAAAAAGCTGCTGAAAAAGCAAAGGAATCTTCTAGTCAAAGTAATCTTCTGGTAG
TGCTTCTAATGAGTCTTATAATGGATCTTCCAATTCAAATGTAGATTATAGTTCATCTGAACAACTAA
TGGATATTCAAATAATTATGGCGGTCAAGATTATCTGGTTCAGGAGATAGTTCAACAAATGGTGGATC
ATCAGAACAAATATTCATCTAGCAATTCAAACAGCGGAGCAAATAATGTCTACAGATATAAAGGCACTGG
TGCTGACGGCTATCAAAGATACTACTACAAAGATCATAATAATGGAGATGTGTATGATGACGATGGAAA
TTACCTTGGAACCTTTGGTGGCGGCATTGCAGAACCTAGTCAACGC

SP082 amino acid (SEQ ID NO:140)

IVQLEKDSKSDKEQVDKLFESFDASSDESISKLELSETSLKTDAGKDYLNKVKESSKAIVDFHLQKG
LAYDVKDSDDKFKDKATLETNVKEITKQIDFIKKVDETFKQENLEETLKSLNDLVDKYQKQIELLKKEE
EKAAEKAAEKAKESSQSNSGASNESYNGSSNSNVYSSSEQTNGYSNNYGGQDYSGSGDSSSTNGGS
SEQYSSNSNSGANNVYRYKGTGADGYQRYYYKDHNNGDVYDDGNYLGNFGGGIAEPSQR

SP083 nucleotide (SEQ ID NO:141)

TCTGACCAAGCAAAAAGAAGCAGTCAATGACAAAGGAAAAGCAGCTGTTGTTAAGGTGGTGGAAAGCCA
GGCAGAACTTTATAGCTTAGAAAAGAATGAAGATGCTAGCCTAAGAAAGTTACAAGCAGATGGACGCAT
CACGGAAGAACAGGCTAAAGCTTATAAAGAATACAAATGATAAAAATGGAGGAGCAAAATCGTAAAGTCAA
TGAT

SP083 amino acid (SEQ ID NO:142)

LTKQKEAVNDKGKAAVVKVVESQAELYSLEKNEDASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVN
D

SP084 nucleotide (SEQ ID NO:143)

GTCCGGCTCTGTCCAGTCCACTTTTTCAGCGGTAGAGGAACAGATTTTCTTTATGGAGTTTGAAGAACT
CTATCGGGAAACCCAAAAACGCAGTGTAGCCAGTCAGCAAAGACTAGTCTGAACTTAGATGGGCAGAC
GCTTAGCAATGGCAGTCAAAGTTGCCAGTCCCTAAAGGAATTCAGGCCCATCAGGCCAAAGTATTAC
ATTTGACCGAGCTGGGGGCAATTCGTCCCTGGCTAAGGTTGAATTTAGACCAGTAAAGGAGCGATTTCG
CTATCAATTATATCTAGGAAATGGAATAATTAACGCATTAAGGAAACAAAAAAT

SP084 amino acid (SEQ ID NO:144)

SGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSNLNDGQTLNNGSQKLPVPKGIQAPSGQSIT
FDRAGGNSSLAKVEFQTSKGAIRYQLYLGNGKIKRIKETKN

SP085 nucleotide (SEQ ID NO:145)

GGGACAAATTCAAAAAAATAGGCAAGAGGAAGCAAAAATCTTGCAAAGGAAGAAGTCTTGAGGGTAGC
TAAGATGGCCCTGCAGACGGGGCAAAATCAGGTAAGCATCAACGGAGTTGAGATTCAGGTATTTCTAG
TGAAAAGGATTGGAGGTCTACCATGGTTCAGAACAGTTGTTGGCAATCAAAGAGCCA

SP085 amino acid (SEQ ID NO:146)

GQIQKNRQEEAKILQKEEVLRVAKMALQTGQNVQSINGVEIQVFSSEKGLEVYHGSEQLLAIKEP

SP086 nucleotide (SEQ ID NO:147)

TCGCTACCAGCAACAAAGCGAGCAAAAGGAGTGGCTCTTGTGTTGTGGACCAACTTGAGGTAGAAATTAGA
CCGTTTCGCAGTTCGAAAAAGTAGAAGGCAATCGCCTATACATGAAGCAAGATGGCAAGGACATCGCCAT
CGGTAAGTCAAAGTCAGATGATTTCGTAACGAATGCTCGTGGTCGAGGTATCAGCCTATGGTTTA
TGGACTCAAATCTGTACGGATTACAGAGGACAATCAACTGGTTCGCTTTCATTTCAGTTCAAAAAGG
CTTAGAAAGGGAGTTCATCTATCGTGTGAAAAAGAAAAAGT

SP086 amino acid (SEQ ID NO:148)

RYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVY
GLKSVRITEDNQLVRHFHFQFQKGLEREFYRVEKEKS

SP087 nucleotide (SEQ ID NO:149)

GAACCGACAAGTCGCCCCACTATCAAGACTATGCTTTGAATAAAGAAAAATTGGTTGCTTTTGCTATGGC
TAAACGAACCAAAGATAAGGTTGAGCAAGAAAGTGGGGAACAGTTTTTTAATCTAGGTCAGGTAAGCTA
TCAAACAAGAAAACTGGCTTAGTGACGAGGGTTCGTACGGATAAGAGCCAATATGAGTTTCTGTTTCC
TTCAGTCAAATCAAAGAAGAGAAAAAGAGATAAAAAGGAAGAGGTAGCGACCGATTCAAGCGAAAAAGT
GGAGAAGAAAAAATCAGAAGAGAAAGCCTGAAAAGAAAGAGAATTCA

SP087 amino acid (SEQ ID NO:150)

NRQVAHYQDYALNKEKLVAFAMAKRTKDKVEQESGEQFFNLGQVSYQNKKTGTVTRVRTDKSQYEFLFP
SVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS

SP088 nucleotide (SEQ ID NO:151)

GGTTGTGCGCTGGCAATATATCCCGTTTCCATCTAAAGGTAGTACAATTGGTCCTTACCCAAATGGTAT
CAGATTAGAAGGTTTTCCAAAGTCAGAGTGGTACTACTTCGATAAAAATGGAGTGCTACAAGAGTTTGT
TGGTTGGAAAACATTAGAGATTAAACTAAAGACAGTGTTGGAAGAAAGTACGGGGAAAAACGTGAAGA
TTCAGAAGATAAAGAAGAGAAGCGTTATTATACGAACATTACTTTAATCAAAATCATTTCTTTAGAGAC
AGGTTGGCTTTATGATCAGTCTAACTGGTATTATCTAGCTAAGACGGAAATTAATGGAGAAAACCTACCT
TGGTGGTGAAAGACGTGCGGGGTGGATAAACGATGATTGCACTTGGTACTACCTAGATCCAACAACCTGG
TATTATGCAAACAGGTTGGCAATATCTAGGTAATAAGTGGTACTACCTCCGTTCCCTCAGGAGCAATGGC
CACTGGCTGGTATCAGGAAGGTACCACTTGGTATTATTTAGACCACCCAAATGGCGATATGAAAACAGG
TTGGCAAAACCTTGGGAACAAATGGTACTATCTCCGTTTCATCAGGAGCTATGGCAACTGGTTGGTATCA
AGATGGTTCAACTTGGTACTACCTAAATGCAGGTAATGGAGACATGAAGACAGGTTGGTTCCAGGTCAA
TGGCAACTGGTACTATGCTTATAGCTCAGGTGCTTTGGCAGTGAATACGACCGTAGATGGCTATTCTGT
CAACTATAATGGCGAATGGGTTCGG

SP088 amino acid (SEQ ID NO:152)

VVGWQYIPFPSKGSTIGPYPNGIRLEGFPKSEWYYFDKNGVLQEFVGWKTLEIKTKDSVGRKYGEKRED
SEMKEEKRYTNYFNFQNHSLGTWLYDQSNWYYLAKTEINGENYLGGERAGWINDDSTWYYLDPTTG
IMQTGWQYLGKWKYYLRSSGAMATGWYQEGTTWYYLDHPNGDMKTGWQNLGNKWYYLRSSGAMATGWYQ
DGSTWYYLNAGNGDMKTGWYFQVNGNWYYAYSSGALAVNTTVDGYSVNYNGEWVR

SP089 nucleotide (SEQ ID NO:153)

GGCCAAATCAGAATGGGTAGAAAGACAAGGGAGCCTTTTATTATCTTGACCAAGATGGAAAGATGAAAAG
AAATGCTTGGGTAGAACTTCTATGTTGGTGCAACAGGTGCCAAAGTAATAGAAGACTGGGTCTATGA
TTCTCAATACGATGCTTGGTTTTATATCAAAGCAGATGGACAGCACGCAGAGAAAGAATGGCTCCAAAT

TAAAGGGAAGGACTATTATTTCAAATCCGGTGGTTATCTACTGACAAGTCAGTGGATTAATCAAGCTTA
TGTGAATGCTAGTGGTGCCAAAGTACAGCAAGGTTGGCTTTTTGACAAACAATACCAATCTTGGTTTTTA
CATCAAAGAAAATGGAACTATGCTGATAAAGAATGGATTTTCGAGAATGGTCACTATTATTATCTAAA
ATCCGGTGGCTACATGGCAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTATCTCAAATTTGA
TGGGAAAATGGCTGAAAAAGAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGG
TGGTTACATGACAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTATCTCAAATCTGATGGGAA
AATAGCTGAAAAAGAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGGTGGTTA
CATGACAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTACCTCAAATCTGATGGGAAAATAGC
TGAAAAAGAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCTGGTGGCTACATGGC
GAAAAATGAGACAGTAGATGGTTATCAGCTTGGGAAGCGATGGTAAATGGCTTGGAGGAAAAACTACAAA
TGAAAAATGCTGCTTACTATCAAGTAGTGCCTGTTACAGCCAATGTTTATGATTGAGATGGTGAAGGCT
TTCCTATATATCGCAAGGTAGTGTGCTATGGCTAGATAAGGATAGAAAAAGTGATGACAAGCGCTTGGC
TATTACTATTTCTGGTTTGTGACGGCTATATGAAAACAGAAGATTTACAAGCGCTAGATGCTAGTAAGGA
CTTTATCCCTTATTATGAGAGTGATGGCCACCGTTTTTATCACTATGTGGCTCAGAATGCTAGTATCCC
AGTAGCTTCTCATCTTTCTGATATGGAAGTAGGCAAGAAATATTATTCGGCAGATGGCCTGCATTTTGA
TGGTTTTAAGCTTGAGAATCCCTTCTTTTCAAAGATTTAAACAGAGGCTACAACTACAGTGTGAAGA
ATTGGATAAGGATTTAGTTTGTCTAAACATTAACAATAGCCTTTTGGAGAACAAGGCGCTACTTTTAA
GGAAGCCGAAGAATACCATATCAATGCTCTTTATCTCCTTGGCCATAGTGCCTTAGAAAGTAAGTAACTG
GGGAAGAAGTAAAATTGCCAAAGATAAGAAATAATTTCTTTGGCATTACAGCCTATGATACGACCCCTTA
CCTTTCTGCTAAGACATTTGATGATGTGGATAAGGGAATTTTAGGTGCAACCAAGTGGATTAAGGAAAA
TTATATCGATAGGGGAAGAACTTTCTTGGAAACAAGGCTTCTGGTATGAATGTGGAATATGCTTCAGA
CCCTTATTGGGGCGAAAAAATTGCTAGTGTGATGATGAAAATCAATGAGAAG

SP089 amino acid (SEQ ID NO:154)

AKSEWVEDKGAFFYYLDQDGKMKRNAWVGTSYVGATGAKVIEDWVYDSQYDAWFYIKADGQHAKEKWLQI
KGKDYFFKSGGYLLTSQWINQAYVNASGAKVQQGWLFQYQSWFYIKENGNADKEWIFENGHYYYLK
SGGYMAANEWIWDKESWFLKFDGKMAKEWVYDSHSQAWYFYSKSGGYMTANEWIWDKESWFLKSDGK
IAKEWVYDSHSQAWYFYSKSGGYMTANEWIWDKESWFLKSDGKIAKEWVYDSHSQAWYFYSKSGGYMA
KNETVDGYQLGSDGKWLGGKTTNENAAYYQVVPVTANVYDSDEKLSYISQGSVWLDKDRKSDDKRLA
ITISGLSGYMKTEDLQALDASKDFIPYYESDGHFRFYHYVAQNASIPVASHLSMEVGGKYYASADGLHFD
GFKLENPFLFKDLTEATNYSAEELDKVFSLLNINNSLLENKGATFKEAEEHYHINALYLLAHSALSNW
GRSKIADKNNFFGITAYDTPYLSAKTFDDVDKGILGATKWIKENYIDRGRFTLGNKASGMNVEYASD
PYWGEKIASVMMKINEK

SP090 nucleotide (SEQ ID NO:155)

ATTTGCAGATGATTCTGAAGGATGGCAGTTTGTCCAAGAAAATGGTAGAACCTACTACAAAAAGGGGA
TCTAAAAGAAACCTACTGGAGAGTGATAGATGGGAAGTACTATTATTTTGATCCTTTATCCGGAGAGAT
GGTTGTGCGCTGGCAATATATACCTGCTCCACACAAGGGGTTACGATTGGTCCTTCTCCAAGAATAGA
GATTGCTCTTAGACCAGATTGGTTTTATTTGGTCAAGATGGTGTATTACAAGAATTTGTTGGCAAGCA
AGTTTTAGAAAGCAAAAACCTGCTACGAATACCAACAAACATCATGGGGAAGAATATGATAGCCAAGCAGA
GAAACGAGTCTATTATTTTGAAGATCAGCGTAGTTATCATACTTTAAAAACTGGTTGGATTTATGAAGA
GGGTCATTGGTATTATTTACAGAAGGATGGTGGCTTTGATTGCGCATCAACAGATTGACGGTTGGAGA
GCTAGCACGTGGTTGGGTTAAGGATTACCTCTTACGTATGATGAAGAGAAGCTAAAAGCAGCTCCATG
GTACTATCTAAATCCAGCAACTGGCATTATGCAAACAGGTTGGCAATATCTAGGTAATAGATGGTACTA
CCTCCATTGCTCAGGAGCTATGGCAACTGGCTGGTATAAGGAAGGCTCAACTTGGTACTATCTAGATGC
TGAAAATGGTGATATGAGAACTGGCTGGCAAAACCTTGGGAACAAATGGTACTATCTCCGTTTCATCAGG
AGCTATGGCAACTGGTTGGTATCAGGAAAGTTCGACTTGGTACTATCTAAATGCAAGTAATGGAGATAT
GAAACAGGCTGGTTCCAAGTCAATGGTAAGTGGTACTATGCCTATGATTGAGGTGCTTTAGCTGTTAA
TACCACAGTAGGTGGTTACTACTTAACTATAATGGTGAATGGGTTAAG

SP090 amino acid (SEQ ID NO:156)

VFADDSEGWQFVQENGRTYYKKGDLKETYWRVIDGKYYFDPPLSGEMVVGWQYIPAPHKGVITIGSPRI
EIALRPDWFYFGQDGVLEQFVGKQVLEAKTATNTNKHGEEYDSQAEKRVYFEDQRSYHTLKTGWIYE
EGHWYYLQKDGDFSRINRLTVGELARGWVKDYPLTYDEEKLKAAPWYYLNPATGIMQTGWQYLGNRWY
YLHSSGAMATGWYKEGSTWYYLDAENGDMRTGWQNLGNKWWYYLRSSGAMATGWYQESSTWYYLNASNGD
MKTGWQFQVNGNWYYAYDSGALAVNTTVGGYYLNYNGEWWK

SP091 nucleotide (SEQ ID NO:157)

TGTCGCTGCAAAATGAAACTGAAGTAGCAAAAACCTTCGCAGGATACAACGACAGCTTCAAGTAGTTCAGAGCAAAATCAGTCTTCTAATAAAACGCAACGAGCGCAGAAGTACAGACTAATGCTGCTGCCACTGGGA
TGGGGATTATTATGTAAAGGATGATGGTTCCTAAAGCTCAAAGTGAATGGATTTTTTGACAACTACTATAA
GGCTTGGTTTTATATTAATTCAGATGGTCGTTACTCGCAGAATGAATGGCATGGAAATTACTACCTGAA
ATCAGGTGGATATATGGCCCAAAACGAGTGGATCTATGACAGTAATTACAAGAGTTGGTTTTATCTCAA
GTCAGATGGGGCTTATGCTCATCAAGAATGGCAATTGATTGGAAATAAGTGGTACTACTTCAAGAAGTG
GGGTTACATGGCTAAAAGCCAATGGCAAGGAAGTTATTTCTTGAATGGTCAAGGAGCTATGATGCAAAA
TGAATGGCTSCTATGATCCAGCCTATTCTGCTTATTTTTATCTAAAATCCGATGGAACCTTATGCTAACC
AAGAGTGGCAAAAAGTGGGCGGCAAAATGGTACTATTTCAAGAAGTGGGGCTATATGGCTCGGAATGAGT
GGCAAGGCAACTACTATTTGACTGGAAGTGGTGCCATGGCGACTGACGAAGTGATTATGGATGGTACTC
GCTATATCTTTGCGGCCTCTGGTGAGCTCAAAGAAAAAAGATTTGAATGTGGGCTGGGTTACAGAG
ATGGTAAGCGCTATTTCTTTAATAATAGAGAAGAACAAGTGGGAACCGAACATGCTAAGAAAGTCATTG
ATATTAGTGAGCACAATGGTCGTATCAATGATTGGAAAAAGGTTATTGATGAGAACGAAGTGGATGGTG
TCATTGTTCTGCTAGGTTATAGCGGTAAAGAAGACAAGGAATTGGCGCATAACATTAAGGAGTTAAACC
GTCTGGGAATTTCCTTATGGTGTCTATCTCTATACCTATGCTGAAAATGAGACCGATGCTGAGAGTGACG
CTAAACAGACCATTGAACTTATAAAGAAATACAATATGAACCTGTCTTACCCTATCTATTATGATGTTG
AGAATTGGGAATTGTAATAAGAGCAAGAGAGCTCCAAGTGATACAGGCACCTGGGTTAAAAATCATCA
ACAAGTACATGGACACGATGAAGCAGGCGGTTATCAAATGTGTATGTCTATAGCTATCGTAGTTTAT
TACAGACGCGTTTAAAAACCCAGATATTTTAAACATGTAACTGGGTAGCGGCCTATACGAATGCTT
TAGAATGGGAAAACCTCATTATTCAGGAAAAAAGGTTGGCAATATACCTCTTCTGAATACATGAAAG
GAATCCAAGGGCGCGTAGATGTCAGCGTTTGGTAT

SP091 amino acid (SEQ ID NO:158)

VAANETEVAKTSQDTTSSSSSEQNQSSNKTQTSAEVQTNAAAHWDGDYVVKDDGSKAQSEWIFDNYYK
AWFYINSDGRYSQNEWHGNYLKSQGYMAQNEWIYDSNYKSWFYLKSDGAYAHQEWQLIGNKWYFCKW
GYMAKSQWQGSYFLNGQGAMQNEWLYDPAYSAYFYLKSDGTYANQEWQKVGGKWYFCKWGYMARNEW
QGNYYLTGSGAMATDEVIMDGTTRYIFAASGELKEKKDLNVGWVHRDGRYFFNNREEQVGTETHAKKVID
ISEHNGRINDWKVIDENEVDGVIVRLGYSGKEDKELAHNIKELNRLGIPYGVYLYTYAENETDAESDA
KQTIELIKKYNMNLSPYIYDVENWEYVNSKRAPSDTGTWVKIINKYMDTMKQAGYQNVVYSYRSL
QTRLKHPDILKHVNWVAAYTNALEWENPHYSKKGWQYTSSEYMKGIQGRVDVSVWY

SP092 nucleotide (SEQ ID NO:159)

TACGTCTCAGCCTACTTTTGTAAAGAGCAGAAGAATCTCCACAAGTTGTGCAAAAATCTTTCATTAGAGAA
GAAATATGAGGAAGCAAAAGCAAAAGCTGATACTGCCAAGAAAGATTACGAAACGGCTAAAAAGAAAGC
AGAAGACGCTCAGAAAAAGTATGAAGATGATCAGAAGAGAAGCTGAGGAGAAAGCTCGAAAAAGAGCAGA
AGCATCTCAAAAATTGAATGATGTGGCGCTTGTGTTCAAAATGCATATAAAGAGTACCGAGAAGTTCA
AAATCAACGTAGTAAATATAAATCTGACGCTGAATATCAGAAAAAATTAACAGAGGTGACTCTAAAAT
AGAGAAGGCTAGGAAAGAGCAACAGGACTTGCAAAATAAATTTAATGAAGTAAGAGCAGTTGTAGTTCC
TGAACCAAATGCGTTGGCTGAGACTAAGAAAAAAGCAGAAGAAGCTAAAGCAGAAGAAAAAGTAGCTAA
GAGAAAATATGATTATGCAACTCTAAAGGTAGCACTAGCGAAGAAAGAAGTAGAGGCTAAGGAACCTGA
AATTGAAAAAATTCAATATGAAATTTCTACTTTGGAACAAGAAGTTGCTACTGCTCAACATCAAGTAGA
TAATTTGAAAAAATTCTTGCTGGTGCGGATCCTGATGATGGCACAGAAGTTATAGAAGCTAAATTTAA
AAAAGGAGAAGCTGAGCTAAACGCTAAACAAGCTGAGTTAGCAAAAAACAAACAGAAGCTTGAAAACT
TCTTGACAGCCTTGATCCTGAAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTT
GGATAAAAAAGCTGATGAACTTCAAAAATAAAGTTGCTGATTTAGAAAAAGAAATTAGTAACCTTGAAAT
ATTACTTGGAGGGGCTGATNCTGAAGATGATACTGCTGCTCTTCAAAATAAATTAGCTACTAAAAAAGC
TGAATTGGAAAAAATCAAAAAGAATTAGATGCAGCTCTTAATGAGTTAGGCCCTGATGGAGATGAAGA
AGAACTCCAGCGCCGGCTCCTCAACCAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTCC
AGCTCCAAAACCAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTCCAGCTCCAAAACCAGA
GCAACAGAGCTCCAGCTCCAAAACCAGAGCAACCAGCTAAGCCGGAGAAACCAGCTGAAGAGCCTACTCA
ACCAGAAAAACAGCCACTCCAAAACAGGCTGGAAACAAGAAACCGGTATGTGGTATTCTACAATAC
TGATGGTTCAATGGCAATAGGTTGGCTCCAAAACAACGGTTTCATGGTACTACCTAAACGCTAACGGCGC
TATGGCAACAGGTTGGGTGAAAGATGGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAAAGC
AAGCCAATGGTTCAAAGTATCAGATAAAATGGTACTATGTCAACAGCAATGGCGCTATGGCGACAGGCTG
GCTCCAATACAATGGCTCATGGTACTACCTCAACGCTAATGGTGATATGGCGACAGGATGGGCTCCAATA
CAACGGTTTCATGGTATTACCTCAACGCTAATGGTGATATGGCGACAGGATGGGCTAAAGTCAACGGTTC
ATGGTACTACCTAAACGCTAACGGTGCTATGGCTACAGGTTGGGCTAAAGTCAACGGTTCATGGTACTA

CCTAAACGCTAACGGTTCAATGGCAACAGGTTGGGTGAAAGATGGAGATACCTGGTACTATCTTGAAGC
ATCAGGTGCTATGAAAGCAAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATGTCAATGGCTTAGG
TGCCCTTGCAGTCAACACAACCTGTAGATGGCTATAAAGTCAATGCCAATGGTGAATGGGTT

SP092 amino acid (SEQ ID NO:160)

TSQPTFVRAEESPOVVEKSSLEKKYEEAKAKADTAKKDYETAKKKAEDAQKKYEDDQKRTEEKARKEAE
ASQKLNDVALVVQNAYKEYREVQNQRSKYKSDAEYQKKLTEVDSKIEKARKEQQDLQNKFNVRVAVVP
EPNALAETKKKAEEAKAEKVAKRKYDYATLKVALAKKEVEAKELEIEKLQYEISTLEQEVATAQHQVD
NLKKLLAGADPDDGTEVIEAKLKKGEAELNAKQAEAKKQTELEKLLDSLDPGKTQDELDKEAEAEEL
DKKADELQNKVADLEKEISNLEILLGGADXEDDTAALQNKLATKKALEKTQKELDAALNELGPDGDEE
ETPAPAPQPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAPAPKPEQPAKPEKPAEEPTQ
PEKPATPKTGWKQENGWYFYNTDGSMAIGWLQNGSWYYLNANGAMATGWVKDGDWYYLEASGAMKA
SQWFKVSDKWYYVNSNGAMATGWLQYNGSWYYLNANGDMATGWLQYNGSWYYLNANGDMATGWAKVNGS
WYYLNANGAMATGWAKVNGSWYYLNANGSMATGWVKDGDWYYLEASGAMKASQWFKVSDKWYYVNLGL
ALAVNTTVDGYKVNANGWV

P093 nucleotide (SEQ ID NO:161)

TGGACAGGTGAAAGGTCATGCTACATTTGTGAAATCCATGACAACCTGAAATGTACCAAGAACAACAGAA
CCATTCTCTCGCCTACAATCAACGCTTGGNTTCGCAAAATCGCATTGTAGATCCTTTTTTGGCGGAGGG
ATATGAGGTCAATTACCAAGTGTCTGACGACCTGATGCAGTCTATGGTTACTTGTCTATTCCAAGTTT
GGAAATCATGGAGCCGGTTTATTTGGGAGCAGATTATCATCATTTAGGGATGGGCTTGGCTCATGTGGA
TGGTACACCGCTGCCTCTGGATGGTACAGGGATTTCGCTCAGTGATTGCTGGGCACCGTGCAGAGCCAAG
CCATGTCTTTTTCCGCCATTTGGATCAGCTAAAAGTTGGAGATGCTCTTTATTATGATAATGGCCAGGA
AATTGTAGAATATCAGATGATGGACACAGAGATTATTTTACCGTCGGAATGGGAAAAATTAGAATCGGT
TAGCTCTAAAAATATCATGACCTTGATAACCTGCGATCCGATTCTACCTTTAATAAACGCTTATTAGT
GAATTTTGAACGAGTGCCTGTTTATCAAAAATCAGATCCACAAACAGCTGCAGTTGCGAGGGTTGCTTT
TACGAAAGAAGGACAATCTGTATCGCGTGTGCAACCTCTCAATGGTTG

SP093 amino acid (SEQ ID NO:162)

GQVKGHATFVKSMTTEMYQEQQNHSLAYNQRLXSQNRIVDPFLAEGYEVNYQVSDDPDAVYGYLSIPSL
EIMEPVYLGADYHHLGMLAHVDGTPLPLDGTGIRSVIAGHRAEP SHVFFRHLQDLKVGDALYYDNGQE
IVEYQMMDEIILPSEWEKLESVSSKNIMTLITCDPIPTFNKRLLVNFERVAVYQKSDPQTAAVARVAF
TKEGQSVSRVATSQWL

SP094 nucleotide (SEQ ID NO:163)

GATTGCTCCTTTGAAGGATTTGAGAGAAACCATGTTGGAAATTGCTTCTGGTGCTCAAAATCTTCGTGC
CAAGGAAGTTGGTGCCTATGAACTGAGAGAAAGTAACCTCGCCAATTTAATGCTATGTTGGATCAGATTGA
TCAGTTGATGGTAGCTATTTCGTAGCCAGGAAGAAACGACCCGTCAGTACCAACTTCAAGCCCTTTCGAG
CCAGATTAATCCACATTTCTCTATAACACTTTGGACACCATCATCTGGATGGCTGAATTTTCATGATAG
TCAGCGAGTGGTGCAGGTGACCAAGTCCTTGGCAACCTATTTCCGCTTGGCGCTCAATCAAGGCAAGGA
CTTGATTTGTCTCTCTGACGAAATCAATCATGTCCGCCAGTATCTCTTTATCCAGAAACAACGCTATGG
AGATAAGCTGGAATACGAAATTAATGAAAATGTTGCCTTTGATAATTTAGTCTTACCCAAGCTGGTCTCT
ACAACCCCTTGTAGAAAATGCTCTTTACCATGGCATTAAGGAAAAGGAAGGTCAGGGCCATATTAACCT
TTCTGTCCAGAAACAGGATTCGGGATTGGTTCATCCGTATTGAGGATGATGGCGTTGGCTTCCAAGATGC
TGGTGATAGTAGTCAAAGTCAACTCAAACGTGGGGGAGTTGGTCTTCAAATGTGATCAACGGCTCAA
ACTTCATTTTGGAGCCAATTACCATATGAAGATTGATTCTAGACCCCAAAAAGGGACGAAAGTTGAAAT
ATATATAAATAGAATAGAACTAGC

SP094 amino acid (SEQ ID NO:164)

IAPLKDLRETMLEIASGAQNLRKEVGAYELREVTRQFNAMLDQIDQLMVAIRSQEETTRQYQLQALSS
QINPHFLYNTLDTIIWMAEFHDSQRVVQVTKSLATYFRLALNQKDLICLSDEINHVRQYLFIQKQRYG
DKLEYEINENVAFDNLVLPKLVLPVENALYHGIKEKEGQGHIKLSVQKQDSGLVIRIEDDGVGFQDA
GDSSQSQLKRGVGLQNVLDQRLKLHFGANYHMKIDSRPQKGTKEIYINRIETS

SP095 nucleotide (SEQ ID NO:165)

TAGGTCATATGGGACTTTTTTCTACAACAAAATAGGCTCCATAATATCTATAAGGGATTTACCCACTA
CAAATATTATAGAGCCGAAAATTCACATCTAATATATGCAGACTACTTTGAAATGAAATTAATAAATTT
ATTAAAGGATGACACAAAAGTTTTTGAATAATCTACATTCAAATTTGTAGAAGGATATAAATAATACCT

GACAGAATCTAAAGAATCTGGAATTAAACAAATGGACAATGTCATAAAATATTTTGAGTTTATTGAATC
TAAAAGTATTGCTTTATATTTTCAAAAACGATTAAATGAGCTGATAGAT

SP095 amino acid (SEQ ID NO:166)

RSYGTFFLQONRLHNIYKGFTHYKYRAENSHLIYADYFEMKLLKLLKDDTKVFEKSTFKFVEGYKIYL
TESKESGIKQMDNVIKYFEFIESKSIALYFQKRLNELID

SP096 nucleotide (SEQ ID NO:167)

CAACGTTGAGAATTATTTGCGAATGTGTTTGGATAGCATTGAGAATCAGACGTATCAAAATTTTGAGTG
TTTATTAATCAATGATGGCTCTCCAGATCATTTCATCCAAAATATGTGAAGAATTTGTAGAGAAAGATTC
TCGTTTCAAAATATTTTGAGAAAGCAAACGCGGCTTTTCATCAGCTCGTAACCTAGGTATTGAATGTTT
GGGGGGGGCGTACATTACTTTTGTAGACTC

SP096 amino acid (SEQ ID NO:168)

NVENYLRMCLDSIQNTYQNFECCLINDGSPDHSSKICEEFVEKDSRFKYFEKANGGLSSARNLGIECS
GGGVHYFCRL

SP097 nucleotide (SEQ ID NO:169)

CTACTATCAATCAAGTTCTTCAGCCATTGAGGCCACCATTTGAGGGCAACAGCCAAACGACCATCAGCCA
GACTAGCCACTTTTATTCAGTCTTATATCAAAAACTAGAAACCACCTCGACTGGTTTGACCCAGCAGAC
GGATGTTCTGGCCTATGCTGAGAATCCCAGTCAAGACAAGGTCGAGGGAATCCGAGATTTGTTTTGAC
CATCTTGAAGTCAGATAAGGACTTGAAAACGTGTGTGCTGGTGACCAAATCTGGTCAGGTCATTTCTAC
AGATGACAGTGTGCAGATGAAAACCTCCTCTGATATGATGGCTGAGGATTGGTACCAAAGGCCATTCA
TCAGGGAGCTATGCCTGTTTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGGTCATTTCTGTCACTCA
AGAACTTGTTGATGCAAAGGGAGCCAATCTTGGTGTGCTTCGTTTGGATATTTCTTATGAAACTCTGGA
AGCCTATCTCAATCAACTCCAGTTGGGGCAGCAGGGCTTTGCCTTCATTATCAATGAAAACCATGAATT
TGTCTACCATCTCAACACACAGTTTATAGTTCTGCTAGCAAAATGGAGGCTATGAAACCTACATCGA
TACAGGTCAGGGTTATACTCCTGGTCACAAATCCTACGTCAGTCAAGAGAAGATTGCAGGAACGTATTG
GACGGTGCTTGGCGTGTATCATTGGAAGTTAGACCAGGTTTCGGAGTCAG

SP097 amino acid (SEQ ID NO:170)

YYQSSSSAIEATIEGNSQTTISQTSHFISYIKKLETTSTGLTQQTDLVLAENPSSQDKVEGIRDLEFLT
ILKSDKDLKTIVLVTKSGQVISTDDSVQMKTSDDMAEDWYQKAIHQGAMPVLT PARKSDSQWVISVTQ
ELVDKAGANLGVLRLDISYETLEAYLNQLQLGQQGFAFIINENHEFVYHPQHTVYSSSSKMEAMKPYID
TGQGYTPGHKSYVSQEKIAGTDWTVLGVSSLEKLDQVRSQ

SP098 nucleotide (SEQ ID NO:171)

GACAAAAACATTAAACGTCCTGAGGTTTTATCACCTGCAGGGACTTTAGAGAAGCTAAAGGTAGCTGT
TCAGTATGGAGCAGATGCTGTCTTTATCGGTGGTCAGGCCTATGGTCTTCGTAGCCGTGCGGGAAACTT
TACTTTTCAACAGATGGAAGAAGGCGTGCAGTTTGCAGGCAAGTATGGTGCCAAGGTCTATGTAGCGGC
TAATATGGTTATGCACGAAGGAAATGAAGCTGGTGTGGTGAGTGGTTCCGTAAACTGCGTGATATCGG
GATTGCAGCAGTTATCGTATCTGACCCAGCCTTGATTATGATTGCAGTGAAGCACCAGGCCTTGA
AATCCACCTTTCTACCCAAGCCAGTGCCACTAACTATGAAACCCCTTGAGTTCTGGAAAGAGCTAGGCTT
GACTCGTGTGCTTTTAGCGCGTGAGGTTTCAATGGAAGAATTAGCTGAGATCCGCAAAACGTACAGATGT
TGAAATTGAAGCCTTTGTCCATGGAGCTATGTGTATTTTCACTCTGGACGTTGTACTCTTTCAAACCA
CATGAGTATGCGTGATGCCAACCGTGGTGGATGTTCTCAGTCATGCCGTTGGAAATACGACCTTTACGA
TATGCCATTTGGGAAAGAACGTAAGAGTTTGCAGGGTGAGATTCCAGAAGAATTTTCAATGTCAGCCGT
TGACATGTCTATGATTGACCANATTCCAGATATGATTGAAAATGGTGTGGACAGTCTAAAAATCGAAGG
ACGTATGNAGTCTATTCACTANGTATCAACAGTAACCAACTGCTACAAGGCGGCTGTGGATGCCTATCT
TGAAAGTCTTGAAAAGTTTGAAGCTATCAAACAAGACTTGGTGGACGAGATGTGGAAGGTTGCCCAACG
TGAACCTGGCTACAGGATTTTACTATGGTACACCATCTGAAAATGAGCAGTTGTTTGGTGTCTCGTCTGTA
AATCCCTGAGTACAAGTTTGTGCTGAAGTTTCTTATGATGATGCGGCACAAACAGCAACTATTTCG
TCAACGAAACGTCATTAACGAAGGGGACCAAGTTGAGTTTTATGGTCCAGGTTTCCGTCAATTTTGAAC
CTATATTGAAGATTTGCATGATGCTAAAGGCAATAAAATCGACCGCGCTCCAAATCCAATGGAACATT
GACTATTAAAGTCCCAACCTGTTCAATCAGGAGACATGGTTCGAGCTCTTAAAGAGGGGCTTATCAA
TCTTTATAAGGAAGATGGAACAGCGTCACAGTTCGTGCT

SP098 amino acid (SEQ ID NO:172)

TKTLKRPEVLSPAGTLEKLKVAVQYGADAVFIGGQAYGLRSRAGNFTFEQMEEGVQFAAKYGAKVYVAA
NMVMHEGNEAGAGWEFRKLRLDIGIAAVIVSDPALIMIAVTEAPGLEIHLSTQASATNYETLEFWKELGL
TRVVLAREVSMEEELAEIRKRTDVEIEAFVHGAMCISYSGRCTLNHNMSMRDANRGGCSQSCRWKYDLYD
MPFGKERKSLQGEIPEEFSSMSAVDMSMIDXIPDMIENGVDLSLKIEGRMXSIHXVSTVTNCKYAAVDAYL
ESPEKF EAIKQDLVDEMWKVAQRELATGFYYGTPSENEQLFGARRKIPEYKFVAEVVSYDDAAQTATIR
QRNVINEGDQVEFYGPGRHFETYIEDLHDAKGNKIDRAPNPMELLTIKVPQPVQSGDMVRALKEGLIN
LYKEDGTSVTVRA

SP099 nucleotide (SEQ ID NO:173)

TTCTCAGGAGACCTTTAAAAATATCACCAATAGCTTCTCCATGCAAATCAATCGTCGCGTCAACCAAGG
AACGCCCTCGTGGTGTCTGGGAATATCAAGGGTGAAGACATCAAAAAATCACCGAAAAACAAGGCCATTGA
GTCTTATGTCAAACGTATCAACGCTATCGGAGATTGACTGGATATGACCTGATTGAAACGCCAGAAAC
CAAGAAGAATCTCACTGCTGATCGTGCCAAAGCGTTTGGAAAGTAGCTTGATGATTACAGGTGTCAATGA
CTCCTCTAAAGAAGACAAGTTTGTCTCTGGTTCTTATAAACTAGTCGAAGGAGAGCACTTAACCAACGA
CGACAAGGATAAAATCCTCTTGCACAAGGACTTGGCAGCCAAACACGGCTGGAAAGTAGGGGACAAGGT
TAAACTGGACTCTAATATCTACGATGCAGATAATGAAAAAGGAGCCAAGGAAACAGTTGAAGTGACAAT
CAAGGGACTCTTTGATGGTCATAAAGTCAGCAGTAACCTACTCACAAGAACTTTACGAAAAACACAGC
TATTACAGACATTACACTGCTGCAAAAACCTTTATGGATACACAGAAGACACAGCCATTTATGGGGACGC
AACCTTCTTTGTAACAGCAGACAAGAACTTGGATGATGTTATGAAAGAGTTGAATGGCATCAGTGGTAT
CAACTGGAAGAGCTACACACTCGTCAAGAGCTCCTCTAACTACCCAGCTCTTGAGCAATCTATCTCTGG
TATGTACAAGATGGCCAAC

SP099 amino acid (SEQ ID NO:174)

SQETFKNITNSFSMQINRRVNQGTPRGAGNIKGEDIKKITENKAIESYVKRINAIGDLTGVDLIETPET
KKNLTADRAKRFGSSLMITGVNDSSKEDKFVSGSYKLVEGEHLTNDKDKILLHKDLAAKHGWKVGDKV
KLDSNIYDADNEKGAKETVEVTIKGLFDGHNKSAVTYSQELYENTAITDIHTAAKLYGYTEDTAIYGDA
TFFVTADKNLDDVMKELNGISGINWKSXYTLVKSSSNYPALQSIISGMYKMAN

SP100 nucleotide (SEQ ID NO:175)

AGTAAATGCGCAATCAAATTCATTAATATTAATAGATGAACCTGAAATCTCACTTCATCCGAGTGCAAT
CTATAAATTTAAAGAGTTTTACTTCAAGAGTGTTTAAATAAAAAACATCAAATTATTATCACTACACA
TTCTACAAACTTATAAAAGATTTTCCTAGAGAAGCCGTGAAACTTTTAGTGAAAAACGGAGAAAAAGT
AGATGTTTATTGAAAATATTGATTATCAGGATGCATTTTGAATTAGGTGATGTGATCATCTTAGGAA
GATGATTTATGTTGAAGATAGACTAGCTAAATATATTCTAGAGTTTGTTATCACTCATTCAGGTAGTGA
GAATCTTAAACAGAATTTAGTAGTGAGATATATTCCTGGTGGAGCAAATCAAATAATTTGTAATAATAT
TTTAAACTCATCGTATTTAGATTCCGATAACCATTAATTTTGGCTTGATGGAGATCAAAACACTAATGT
TAGTGAATCAAATAATTTAATGAACTATCTTGAAAATGGTGTGTTATATCAGATAAAATTCCTGAATC
AGATAATAAAATCTTGATGATATTATAAAATTGATAANGGGATGTCCAATTAAATTTAATGTTTCAGG
TAATAAAGGGCAAAAAATAATATTGAATTAATTGCGAAACAAAGAAGCTTTATAGATTATTGGGCTAA
ATAC

SP100 amino acid (SEQ ID NO:176)

VNAQNSNLILIDEPEISLHPSAIYKFKEFLQECNLKKHQIIITTHSTQLIKDFPREAVKLLVKNGEKV
DVNIENIDYQDAFFELGDVYHSRKMIYVEDRLAKYILEFVITHSGSENKQNLVVRYIPGGANQIIICNNI
LNSSYLDSDNHYFWLDGDQNTNVSESNLMNYLENGVVISDKIPESDNKNLDDIIKLI XGCPKIFNVSG
NKGQKNNIELIAKQRSFIDYWAKY

SP101 nucleotide (SEQ ID NO:177)

TTACCGCGTTCATCAAGATGTCAAACAAGTCATGACCTATCAACCCATGGTGGCAGAAAATATTGAGTGA
ACAAGACACCCCAGCAAACGAAGAGCTGTGCTTGCTATGATTTATACTGAAACAAAAGGAAAAGAAGG
CGATGTTATGCAGTCTAGTGAGTCTGCAAGTGGTTCCACCAACACCATCAATGATAATGCCTCTAGCAT
TCGGCAAGGCATTCAAACCTCTGACAGGCAATCTCTATCTGGCGCAGAAGAAGGGGGTAGATATCTGGAC
AGCTGTTCAAGCCTATAATTTTGGACCTGCCTATATCGATTTTATCGCCCAAAATGGCAAGGAAAATAC
CCTGGCTCTAGCCAAACAGTACTCTCGTGAGACTGTTGCCCCCTTGCTTGGTAATAGGACTGGAAAGAC
TTATAGTTATATTCACCCCATTTCCATTTTTCACGGTGCTGAACTCTATGTAAATGGAGGAAACTATTA
TTATTCTAGACAGGTACGACTTAACCTTTACATCATCAAATGTTTCACTCTCTTTTCAACATCTGGC

SP101 amino acid (SEQ ID NO:178)

YRVHQDVKQVMYQPMVREILSEQDTPANEELVLAMIYTETKGKEGDMQSSSESASGSTNTINDNASSI
RQGIQTLTGNLYLAQKKGVDIWTAVQAYNFGPAYIDFIAQNGKENTLALAKQYSRETVPALLGNRTGKT
YSYIHPISIFHGAELYVNGGNYYSRQVRLNLYIIKCFTLFSTSG

SP102 nucleotide (SEQ ID NO:179)

GTGGATGGGCTTTAACTATCTTCGTATTCGCCGTGCGGCTAAAATTGTGGACAATGAGGAGTTTGAAGC
CTTGATTTCGTACGGGTCAATTGATTGATTTCGCGGACCCAGCAGAATTCCACAGAAAACATATCCTTGG
TGCACGCAATATTTCCTTCAAGTCAGTTGAAAAC TAGTCTTGACGCCCTTCGTAAAGATAAACCTGTCCT
TCTCTACGAAAACCAACGTGCGCAACGAGTTACAAATGCAGCTCTTTACTTGAAAAACAAGGTTTTTC
TGAGATTTATATCCTTTCTTATGGCTTGGATTCTTGGAAGGGAAAGTGAAGACTAGC

SP102 amino acid (SEQ ID NO:180)

WMGFNYLRIRRAAKIVDNEEFELIRTGQLIDLRDPAEFHRKHILGARNIPSSQLKTSLAALRKDKPVL
LYENQRAQRVNTAALYLKKQGFSEIYILSYGLDSWKGVKTS

SP103 nucleotide (SEQ ID NO:181)

ACTAAACCAGCATCGTTTCGAGGAAAATAAGGACAATAATCGTGTCTCTTATGTGGATGGCAGCCAGTC
AAGTCAGAAAAGTGAAAACCTTGACACCAGACCAGGTTAGCCAGAAAAGAAGGAATTCAGGCTGAGCAAAT
TGTAATCAAAATTACAGATCAGGGCTATGTAACGTCACACGGTGACCACTATCATTACTATAATGGGAA
AGTTCCTTATGATGCCCTCTTTAGTGAAGAACCTTGATGAAGGATCCAACTATCAACTTAAAGACGC
TGATATTGTCAATGAAGTCAAGGGTGGTTATATCATCAAGGTCGATGGAAAATATTATGTCTACCTGAA
AGATGCAGCTCATGCTGATAATGTTTGAACATAAGATGAAATCAATCGTCAAAAACAAGAACATGTCAA
AGATAATGAGAAGGTTAACTCTAATGTTGCTGTAGCAAGGCTCAGGGACGATATACGACAAATGATGG
TTATGTCTTTAATCCAGCTGATATTATCGAAGATACGGGTAATGCTTATATCGTTCCTCATGGAGGTCA
CTATCACTACATTCCCAAAAGCGATTTATCTGCTAGTGAATTAGCAGCAGCTAAAGCACATCTGGCTGG
AAAAAATATGCAACCGAGTCAGTTAAGCTATTCTTCAACAGCTAGTGACAATAACACGCAATCTGTAGC
AAAAGGATCAACTAGCAAGCCAGCAAAATAAATCTGAAAATCTCCAGAGTCTTTTGAAGGAACTCTATGA
TTCACCTAGCGCCCAACGTTACAGTGAATCAGATGGCCTGGTCTTTGACCCTGCTAAGATTATCAGTCG
TACACCAAATGGAGTTGCGATTCCGCATGGCGACCATTTACCACTTTATTCCTTACAGCAAGCTTTCTGC
CTTAGAAGAAAAGATTGCCAGAATGGTGCCATCAGTGGAACCTGGTCTTACAGTTTCTACAAATGCAAA
ACCTAATGAAGTAGTGTCTAGTCTAGGCAGTCTTTCAAGCAATCCTTCTTCTTTAACGACAAGTAAGGA
GCTCTCTTTCAGCATCTGATGGTTATATTTTAAATCCAAAAGATATCGTTGAAGAAACGGCTACAGCTTA
TATTGTGAAGCATGGTGTATTTCCATTACATTTCAAAATCAAATCAAATTTGGGCAACCGACTCTTCC
AAACAATAGTCTAGCAACACCTTCTCCATCTCTTCCAATCAATCCAGGAACCTCACATGAGAAACATGA
AGAAGATGGATACGGATTTGATGCTAATCGTATTATCGCTGAAGATGAATCAGGTTTTGTGATGAGTCA
CGGAGACCACAATCATTATTTCTTCAAGAAG

SP103 amino acid (SEQ ID NO:182)

LNQHRSEQENKDNRRVSYVDGSQSSQKSENLTDPQVSQKEGIQAEQIVIKITDQGYVTSHGDHYHYNGK
VPYDALFSEELLMKDPNYQLKDADIVNEVKGYYIIKVDGKYYVYLKDAHADNVRTKDEINRQKQEHVK
DNEKVNNSNAVARSQGRYTTNDGYVFNPAIIEDTGNAYIVPHGGHYHYIPKSDLASASELAAKAHLAG
KNMQPSQLSYSSASDNNTQSVAKGSTSKPANKSENLSLLKELYDSPSAQRYSES DGLVFDPAKIIISR
TPNGVAIPHGDHYHFI PYSKLSALEEKIARMVPI SGTGSTVSTNAKPNEVVSSLGSLSSNPSSLTTSKE
LSSASDGYIFNPKDIVEETATAYIVRHGDHFIYIPKSNQIGQPTLPNNSLATPSPSLPINPGTSHEKHE
EDGYGFDANRIIAEDES GFVMSHGDHNHYFFKK

SP105 nucleotide (SEQ ID NO:183)

TGACTACCTTGAAATCCCACCTTACAGCTATCTTGGTGGATTCAACACTAAAGTTCTTCCAACCTCAAT
GATGAACATCATCAACGGTGGTTCTCACTGACGCTCCAATCGCTTTCCAAGAGTTTCATGATCTTGCC
AGTTGGTGCGCCAACATTTAAAGAAGCCCTTCGTTACGGTGCTGAAATCTTCCACGCTCTTAAGAAAAT
CCTTAAATCACGTGGTTTGGAAACTGCCGTAGGTGACGAAGGTGGATTGCTCCTCGTTTGAAGGAAC
TGAAGATGGTGTGAAACTATCCTTGCTGCGATTGAAGCTGCTGGATATGTACCAGGTAAAGACGTATT
TATCGGATTTGACTGTGCTTCATCAGAATTCTACGATAAAGAACGTAAAGTTTACGACTACACTAAATT
TGAAGGTGAAGGTGCTGCTGTTGCTGACATCTGCAGAACAAATCGACTACCTTGAAGAATTGGTTAACAA
ATACCCAATCATCACTATTGAAGATGGTATGGATGAAAACGACTGGGATGGTTGGAAAGCTCTTACTGA
ACGCTCTGGTAAGAAAGTACAACCTGTTGGTGACGACTTCTTCGTAACAAACACTGACTACCTTGCACG

TGGTATCCAAGAAGGTGCTGCTAACTCAATCCTTATCAAAGTTAACCAAATCGGTACTCTTACTGAAAC
TTTTGAAGCTATCGAAATGGCTAAAAGAAGCTGGTTACACTGCTGTTGTATCACACCGTTCAGGTGAAAC
TGAAGATTCAACAATCGCTGATATTGCAGTTGCAACTAACGCAGGACAAATCAAGACTGGTTCACTTTC
ACGTACAGACCGCATCGCTAAATACAACCAATTGCTTCGTATCGAAGACCAACTTGGTGAAGTAGCTGA
ATATCGTGGATTGAAATCATTCTACAACCTTAAAAAA

SP105 amino acid (SEQ ID NO:184)

DYLEIPLYSYLGGFNTKVLPTPMMNIINGGSHSDAPIAFQEFMILPVGAPTFKEALRYGAEIFHALKKI
LKSRGLETAVGDEGGFAPRFEGTEDGVETILAAIEAAGYVPGKDVFIGFDCASSEFYDKERKVDYDTKF
EGEGAAVRTSAEQIDYLEELVNKYPIITIEDGMDENDWDGWKALTERLGKKVQLVGDDFFVTNTDYLAR
GIQEGAANSILIKVNQIGTLTETFEAIEMAKEAGYTAVVSHRSGETEDSTIADIAVATNAGQIKTGSLS
RTDRIAKYNQLLRIEDQLGEVAEYRGLKSFYNLKK

SP106 nucleotide (SEQ ID NO:185)

TCGTATCTTTTTTTGGAGCAATGTTTCGCGTAGAAGGACATTCCATGGATCCGACCCTAGCGGATGGCGA
AATTCCTCTTCGTTGTAAAACACCTTCCTATTGACCGTTTTGATATCGTGGTGGCCCATGAGGAAGATGG
CAATAAGGACATCGTCAAGCGCGTGATTGGAATGCCCTGGCGACACCATTTCGTTACGAAAATGATAAACT
CTACATCAATGACAAAGAAACGGACGAGCCTTATCTAGCAGACTATATCAAACGCTTCAAGGATGACAA
ACTCCAAAGCACTTACTCAGGCAAGGGCTTTGAGGAAATAAAGGAACCTTTCTTTAGAACTATCGCTCA
AAAAGCTCAAGCCTTCACAGTTGATGTCAACTACAACACCAACTTTAGCTTTTACTGTCCAGAAGGAGA
ATACCTTCTCCTCGGAGATGACCGCTTGGTTTCGAGCGACAGCCGCCACGTAGGTACCTTCAAAGCAAA
AGATATCACAGGGGAAGCTAAATTCCGCTTATGGCCAATCACCCGTATCGGAACATTT

SP106 amino acid (SEQ ID NO:186)

RIFWFSNVRVEGHSMDPTLADGEILFVVKHLPIDRFDIVVAHEEDGNKDIVKRVIGMPGDTIRYENDKL
YINDKETDEPYLADYIKRFKDDKLQSTYSGKGFEGNKGTFFRSIAQKAQAFVVDVNYNTNFSFTVPEGE
YLLLGDDRLVSSDSRHVGTFFAKDITGEAKFRLWPITRIGTF

SP107 nucleotide (SEQ ID NO:187)

GGACTCTCTCAAAGATGTGAAAGCAAATGCTAGCGACAGCAAGCCTGCACAGGACAAGAAGGATGCAAA
ACAAGGAACGGAAGATAGTAAGGATTCAGATAAGATGACTGAAACAAACTCAGTTCCGGCAGGAGTGAT
TGTGGTCAGTCTACTTGCCCTCCTAGGCGTGATTGCCCTTCTGGCTGATTTCGCCGTAAGAAAGAGTCAGA
AATCCAGCAATTAAGCACGGAATTGATCAAGGTTCTAGGACAGCTAGATGCAGAAAAAGCGGATAAAAA
AGTCCTTGCCAAAGCCCAAACCTTCTCCAAGAAACCTTGATTTCGTGAAAGAAAAATGGCTCAGC
AGAGACAGAACTAACTAGTAGAGGAGCTTAAAGCAATCCTTGACAAACTCAAG

SP107 amino acid (SEQ ID NO:188)

DSLKDVKANASDSKPAQDKKDAKQGTEDSKDSKMTETNSVPAGVIVVSLALLGVIAFWLIRRKKESE
IQQLSTELIKVLGQLDAEKADKKVLAKAQNLLQETLDFVKEENGSAETETKLVEELKAILDKLK

SP108 nucleotide (SEQ ID NO:189)

CAAGAAATCCTATCATCTCTTCCAGAAGCAAACAGAGACGAGGGGAATTCAGACTCAGTTGATTGAAGA
ATCGCTTAGTCAGCAGACTATAATCCAGTCCTTCAATGCTCAAACAGAATTTATCCAAAGATTGCGTGA
GGCTCATGACAACACTACTCAGGCTATTCTCAGTCAGCCATCTTTTATTCTTCAACGGTCAATCCTTCGAC
TCGCTTTGTAAATGCACTCATTTATGCCCTTTTAGCTGGAGTAGGAGCTTATCGTATCATGATGGGTTTC
AGCCTTGACCGTCGGTCGTTTAGTGACTTTTTTGAACATATGTTTCAGCAATACACCAAGCCCTTTAACGA
TATTTCTTCAGTGCTAGCTGAGTTGCAAAGTGCTCTGGCTTGCGTAGAGCGTATCTATGGAGTCTTAGA
TAGCCCTGAAGTGGCTGAAACAGGTAAGGAAGTCTTGACGACCAGTGACCAAGTTAAGGGAGCTATTTTC
CTTTAAACATGTCTCTTTTGGCTACCATCCTGAAAAAATTTTGATTAAGGACTTGTCTATCGATATTCC
AGCTGGTAGTAAGGTAGCCATCGTTGGTCCGACAGGTGCTGGGAAATCAACTCTTATCAATCTCCTTAT
GCGTTTTTATCCCATTAGCTCGGGAGATATCTTGCTGGATGGGCAATCCATTATGATTATACACGAGT
ATCATTGAGACAGCAGTTTGGTATGGTGCTTCAAGAAACCTGGCTCACACAAGGGACCATTCATGATAA
TATTGCCTTTGGCAATCCTGAAGCCAGTCGAGAGCAAGTAATTGCTGCTGCCAAAGCAGCTAATGCGAGA
CTTTTTTCATCCAACAGTTGCCACAGGGATACGATACCAAGTTGGAATGCTGGAGAATCTCTCTCTGT
CGGCCAAGCTCAGCTCTTGACCATAGCCGAGTCTTTCTGGCTATTCCAAAGATTCTTATCTTAGACGA
GGCAACTTCTTCCATTGATACACGACAGAAGTGCTGGTACAGGATGCCTTTGCAAACTCATGAAGGG
CCGCACAAGTTTCATCATTGCTCACCGTTTGTCAACCATTACAGGATGCGGATTTAATTCTTGCTTAGT

AGATGGTGATATTGTTGAATATGGTAACCATCAAGAACTCATGGATAGAAAGGGTAAGTATTACCAAAT
GCAAAAAGCTGCGGCTTTTAGTTCTGA

A

SP108 amino acid (SEQ ID NO:190)

KKSYHLFQKQTETRGITQQLIEESLSQQTIIQSFNQTEFIQRLREAHDNYSQSAIFYSSSTVNPST
RFVNALIYALLAGVGAYRIMMGSALTVGRLVTFNLVYQQYTKPFNDISSVLAELQSALACVERIYGVLD
SPEVAETGKEVLTTSDQVKGAISFKHVSFGYHPEKILIKDLSIDIPAGSKVAIVGPTGAGKSTLINLLM
RFYPISSGDILLDGQSIYDYTRVSLRQQFGMVLQETWLTQGTIHDNIAFGNPEASREQVIAAAKAAANAD
FFIQQLPQGYDTKLENAGESLSVGQAQLLTIARVFLAIPKILILDEATSSIDTRTEVLVQDAFAKLMKG
RTSFIIAHRLSTIQDADLILVLVDGDIVEYGNHQLMDRKGKYYQMOKAAAFSSE

SP109 nucleotide (SEQ ID NO:191)

ACGAAATGCAGGGCAGACAGATGCCTCGCAAATTGAAAAGGCGGCAGTTAGCCAAGGAGGAAAAGCAGT
GAAAAAACAGAAATTAGTAAAGACGCAGACTTGCACGAAATTTATCTAGCTGGAGGTTGTTTCTGGGG
AGTGGAGGAATATTTCTCACGTGTTCCCGGGGTGACGGATGCCGTTTCAGGCTATGCAAATGGTAGAGG
AGAAACAACCAAGTACGAATTGATTAACCAACAGGTCATGCAGAAACCGTCCATGTCACCTATGATGC
CAAGCAAATTTCTCTCAAGGAAATCCCTGCTTCACTATTTCCGCATTATCAATCCAACGACAAAAATAA
ACAAGGAAATGATGTGGGGACCCAGTACCGTACTGGTGTATTACACAGATGACAAGGATTTGGAAGT
GATTAACCAAGTCTTTGATGAGGTGGCTAAGAAATACGATCAACCTCTAGCAGTTGAAAAGGAAAACCTT
GAAGAATTTTGTGGTGGCTGAGGATTACCATCAAGACTATCTCAAGAAAAATCCAAATGGCTACTGCCA
TATCAATGTTAATCAGGCGGCCTATCCTGTCAATTGATGCCAGCAAATATCCAAAACCAAGTATGAGGA
ATTGAAAAGACCCTGTACCTGAGGAGTATGCAGTTACCCAGGAAAATCAAACAGAACGAGCTTTCTC
AAACCGTTACTGGGATAAATTTGAATCCGGTATCTATGTGGATATAGCAACTGGGGAACCTCTCTTTTC
ATCAAAGACAAATTTGAGTCTGGTGTGGCTGGCCTAGTTTTACCAACCCATCAGTCCAGATGTTGT
CACCTACAAGGAAGATAAGTCTACAATATGACGCGTATGGAAGTGGGAGCCGAGTAGGAGATTCTCA
CCTTGGGCATGTCTTTACGGATGGTCCACAGGACAAGGGCGGCTTACGTTACTGTATCAATAGCCTCTC
TATCCGCTTTATTCCCAAAGACCAAATGGAAGAAAAAGGCTACGCTTATTTACTAGATTATGTTGAT

SP109 amino acid (SEQ ID NO:192)

RNAGQTDASQIEKAAVSQGGKAVKKTEISKDADLHEIYLAGGCFWGVVEEYFSRVPGVTDVSGYANGRG
ETTKYELINQTHAETVHVTYDAKQISLKEILLHYFRIINPTSKNKQGNVDVGTQYRTGVYYTDDKDLEV
INQVFDEVAKKYDQPLAVEKENLKNFVVAEDYHQDYLLKNPNNGYCHINVNQAAYPVIDASKYPKPSDEE
LKKTLSPPEYAVTQENQTERAFSNRYWDFESGIYVDIATGEPLFSSKDKFESGCGWPSFTQPISPDVV
TYKEDKSYNMTRMEVRSRVGDSHLGHVFTDGPQDKGGLRYCINSLSIRFIPKQMEEEKGYAYLLDYVD

SP110 nucleotide (SEQ ID NO:193)

TGTATAGTTTTTAGCGCTTGTCTTCTAATTCTGNTAAAAATGAAGAAAATACTTCTAAAGAGCATGCC
CCTGATAAAATAGTTTTAGATCATGCTTTCGGTCAAACCTATATTAGATAAAAAACCTGAAAGAGTTGCA
ACTATTGCTTGGGGAAATCATGATGTAGCATTAGCTTTAGGAATAGTTCCTGTTGGATTTTCAAAAGCA
AATTACGGTGTAAAGTGTGATAAAGGAGTTTACCATGGACAGAAGAAAAAATCAAAGAATAAATGGT
AAAGCTAACCTATTTGACGATTTGGATGGACTTAACCTTTGAAGCAATATCAAATTCTAAACCAGATGTT
ATCTTAGCAGGTTATTCTGGTATAACTAAAGAAGATTATGACACTCTATCA

SP110 amino acid (SEQ ID NO:194)

CIVFSACSSNSXKNEENTSKEHAPDKIVLDHAFGQTILDKKPERVATIAWGNHDVALALGIVPVGFASKA
NYGVSADKGVLPWTEEKIKELNGKANLFDLDGLNFEAISNSKPDVILAGYSGITKEDYDTLS

SP111 nucleotide (SEQ ID NO:195)

GTGTGTCGAGCATATTCTGAAGCAAACCTATCAAAATATAGAAATTATTTTAGTTGATGACGGTTCTAC
GGATAATTCTGGGGAAATTTGTGATGCTTTTATGATGCAAGATAATCGTGTGCGAGTATTGCATCAAGA
AAATAAGGGGGGGGCGAGCACAAGCTAAAAATATGGGGATTAGTGTAGCTAAGGGAGAGTACATCACGAT
TGTTGATTTCAGATGATATCGTAAAAGAAAATATGATTGAAACTCTTTATCAGCAAGTCCAAGAAAAGGA
TGCAGATGTTGTTATAGGGAATTACTATAATTATGACGAAAGTGACGGGAATTTTTATTTTTATGTAAC
AGGGCAAGATTTTTGCGTCGAAGAATTAGCTATACAAGAAATTATGAACCGTCAAGCAGGAGATTGGAA
ATTCAATAGCTCGGCCCTTTATATTGCCGACATTTAAGTTGATTAAAAAGAATTATTCAATGAAGTTCA
CTTTTCAAATGGTCGCCGCTTTGATGATGAAGCAACTATGCATCGCTTTTATCTTTTAGCCTCTAAAT
CGTCTTTATAAACGATAATCTCTATCTGTATAGAAGACGTTTCAGGAAGCATCATGAGAACGGAATTTGA

TCTTTCCTGGGCAAGAGATATTGTTGAAGTGTTCCTAAGAAAATATCGGATTGTGTCTTGGCTGGTTT
GGATGTCTCCGTTCTGCGTATTCGATTTGTCAATCTTTTAAAGATTATAAGCAAACCTTAGAATACCA
TCAATTAACAGATACTGAGGAATATAAGATATTTGTTTCAGATTAAAGTTGTTTTTGTATGCAGAACA
AAGAAATGGTAAAGT

SP111 amino acid (SEQ ID NO:196)

CVEHILKQTYQNIEIILVDDGSTDNSGEICDAFMMQDNVRVRLHQENKGGAQAQNMGISVAKGEYITI
VDSDDIVKENMIETLYQQVQEKDADVIGNYNYDESDGNFYFYVTGQDFCVEELAIQEIMNRQAGDWK
FNSSAFILPTFKLIKELFNEVHFSNGRRFDDEATMHRFYLLASKIVFINDNLYLYRRRSGSIMRTEFD
LSWARDIVEVFSKKISDCVLAGLDVSVLRIRFVNLLKDYKQTLLEYHQLTDTEYKDICFRLKLFDAEQ
RNGKS

SP0112 nucleotide (SEQ ID NO:197)

GTGTTTGGATAGCATTCAAGATCAGACGTATCAAAATTTGAGTGTATTATTAATCAATGATGGCTCTCC
AGATCATTCATCCAAAATATGTGAAGAATTTGTAGAGAAAGATTCTCGTTTCAAATATTTTGAGAAAGC
AAACGGCGGTCTTTCATCAGCTCGTAACCTAGGTATTGAATGTTTCGGGGGGGGCGTACATTACTTTTGT
AGACTCTGATGATTGGTTGGAACATGATGCTTTAGACCGATTATATGGTGCTTTGAAAAAGGAAAACGC
AGATATTAGTATCGGGCGTTATAATCTTATGATGAAACACGCTATGTGTATATGACTTATGTTACGGA
TCCAGATGATTCTCTAGAAGTGATAGAAGGTAAAGCAATTATGGATAGGGAAGGTCTCGAAGAAGTCAG
AAATGGGAAGTGGACTGTAGCTGTCTTGAAGTTATTCAAGAGAGAGTTACTACAAGATTACCATTTC
TATAGGAAAAATTCAGAGGATACTTACTGGACATGGAAGGTACTTCTAAGAGCTTCGAGGATAGTCTA
TTTGAATCGTTGTGTTTACTGGTACCGTGTGTTGTTTATCTGATACTTTATCGAATACATGGAGTGAAAA
GCGTATGTATGATGAAATTGGGGCTAGGGAAGAAAAGATAGCTATTTTAGCAAGTTCAGACTATGACTT
GACCAATCATATTTTGATTTATAAAAAATAGATTACAAAGAGTGATAGCAAAATTAGAAGAACAAAATAT
GCAGTTCACAGAGATTTACAGAAGAATGATGGAATAATTGTCTTTACTTCCG

SP0112 amino acid (SEQ ID NO:198)

CLDSIQNQTYQNFECILLINDGSPDHSSKICEEFVEKDSRFKYFEKANGGLSSARNLGIECSSGAYITFV
DSDDWLEHDALDRLYGALKKENADISIGRYNSYDETRYVYMTYVTDPPDSLEVIEGKAIMDREGVEEVR
NGNWTVAVLKLFKRELLQDLFPPIGKIAEDTYWTKVLLRASRIVLYLNRVCVYWRVGLSDTLNNTWSEK
RMYDEIGAREEKIAILASSDYDLTNHILYKNRLQRVIAKLEEQNMQFTEIYRRMMEKLSLLP

SP113 nucleotide (SEQ ID NO:199)

GTGCCTAGATAGTATTATTACTCAACATATAAAAAATTTGAGATTGTTGTCGTTAATGATGGTTCTAC
GGATGCTTCAGGTGAAATTTGTAAAGAATTTTCAGAAATGGATCACCGAATTCTCTATATAGAACAAGA
AAATGCTGGTCTTTCTGCCGCACGAAACACCGGTCTGAATAATATGTCCGGAATTTATGTGACCTTTGT
GGACTCGGATGATTGGATTGAGCAAGATTATGTAGAACTCTATATAAAAAAATAGTAGAGTATCAGGC
TGATATTGCAGTTGGTAATTATTATTCTTTCAACGAAAGTGAAGGAATGTTCTACTTTTCATATATTGGG
AGACTCCTATTATGAGAAAGTATATGATAATGTTTCTATCTTTGAGAACTTGATGAACTCAAGAAAT
GAAGAGTTTGTCTTTGATATCTGCTTGGGGTAACTCTATAAGGCAAGATTGTTTGAGCAGTTGCGCTT
TGACATAGGTAAATTAGGAGAAGATGGTTACCTCAATCAAAGGTATATTTATTATCAGAAAAGGTAAT
TTATTTAAATAAAAGTCTTTATGCTTATCGGATTAGAAAAGGTAGTTTATCAAGAGTTTGGACAGAAAA
GTGGATGCACGCTTTAGTTGATGCTATGTCTGAACGTATTACGCTACTAGCTAATATGGGTTATCCTCT
AGAGAAACACTTGGCAGTTTATCGTCAGATGTTGGAAGTCAGTCTCGCCAACGGTCAAGCTAGTGGTTT
ATCTGACACAGCAACGTATAAAGAGTTTGAATGAAACAAAGGCTTTTAAATCAGCTATCGAGACAAGA
GGAAAGTGAAAGAAAGCCATTGTCTCTGCAGCAAACATATGGCTATGTAGACCAAGTTTTAACGACAAT
CAAGTCTATTTGTTATCATAATCGTTCGATTCTGTTTATCTGATTATAGCGATTTTCCAAATGAATG
GATTAAGCAATTAAATAAGCGCTTAGAGAAGTTTGACTCAGAAATTATTAATTGTCTGGGTAACTTCTGA
GCAAAATTCATGTTATAAATCGGATATTAGTTACACAGTCTTTTACGCTATTTTCATAGCTGATTTCTG
GCAAGAAGACAAGGCCCTCTACTTGGACTGTGATCTAGTTGTAACGAAAAATCTGGATGACTTGTGTTGC
TACAGACTTACAAGATTATCCTTTGGCTGCTGTTAGAGATTTTGGGGGCAGAGCTTATTTGGTCAAGA
AATCTTTAATGCCGGTGTCTCTTGGTAAACAATGCTTTTTGGAAAAAAGAGAATATGACCCAAAAAT
AATTGATGTAACCAATGAATGGCATGATAAGGTGGATCAGGCAGATCAGAGCATCTGAATATGCTTTT
TGAACATAAATGGTTGGAATTGGACTTTGATTATAATCATATTGTCATTCATAAACAGTTTGTCTGATTA
TCAATTGCTCTGAGGGTCAGGATTATCCTGCTATTATTCACTATCTTCTCATCGGAAACCGTGGAAGA
TTTGGCGGCCCAAACCTATCGTGAAGTTTGGTGGTACTATCATGGGCTTGAATGGACAGAATTGGGACA
AAACCATCATTTACATCCATTACAAAGATCTCACATCTATCCAATAAAGGAACCTTTCACTTGTCTAAT
CTATACTGCCTCAGACCATATTGAACAAATGAGACATTGGTTCAATCCTTGCCTGATATTCAGTTTAA

GATAGCAGCTAGAGTAATAGTTAGTGATCGATTGGCTCAGATGACAATTTATCCAAACGTGACTATATT
TAACGGAATTCACATATTTGGTAGATGTCGATAATGAATTGGTAGAAACCAGTCAAGTACTTTTAGATAT
TAATCATGGCGAAAAGACAGAAAGAAATTCCTCGATCAATTTGCTAATCTTGGCAAGCCTATCTTATCCTT
TGAAAATACTAAACCTATGAAGTAGGTCAGGAGGCATATGCTGTTGACCAAGTTCAAGCAATGATTGA
AAAATTGAGAGAAATAAGCAA

SP113 amino acid (SEQ ID NO:200)

CLDSIITQTYKNIEIVVNDGSTDASGEICKFSEMDHRILYIEQENAGLSAARNTGLNNMSGNYVTFV
DSDDWIEQDYVETLYKKIVEYQADIAVGNYYSFNESEGMFYFHLGDSYEEKVYDNVSI FENLYETQEM
KSFALISAWGKLYKARLFEQLRFDIGKLGEDGYLNQKVYLLSEKVIYLNKSLYAYRIRKGSLSRVWTEK
WMHALVDAMSERITLLANMGYPLEKHLAVYRQMLEVSLANGQASGLSDTATYKEFEMQORLLNQLSRQE
ESEKKAIVLAANYGYVDQVLTTIKSICYHNRSIRFYLIHSDFPNEWIKQLNKRLEKFDSEI INCRVTSE
QISCYKSDISYTVFLRYFIADVFQEDKALYLDCLDVVTKNLDDLFDLQDYPLAAVRDFGGRAYFGQE
IFNAGVLLVNNAFWKKENMTQKLIDVTNEWHDKVDQADQSILNMLFEHKWLELDFDYNHIVIHKQFADY
QLPEGQDYPALIIHYLSHRKPWKDLAAQTYREVWYYHGLEWTELGNHHLHPLQRSHIYPIKEPFTCLI
YTASDHIEQIETLVQSLPDIQFKIAARVIVSDRLAQMTIYPNVTIFNGIHYLVVDVDELVETSQVLLDI
NHGEKTEEILDQFANLGKPILSFENTKTYEVGQEAYAVDQVQAMIEKLREISK

SP114 nucleotide (SEQ ID NO:201)

CATTCAGAAGCAGACCTATCAAAATCTGGAATTTATCTTGTGATGATGGTGCAACAGATGAAAGTGG
TCGCTTGTGTGATTCAATCGCTGAACAAGATGACAGGGGTGTCAGTGCTTCATAAAAAAGAACGAAGGATT
GTCGCAAGCACGAAATGATGGGATGAAGCAGGCTCACGGGGATTATCTGATTTTTATTGACTCAGATGA
TTATATCCATCCAGAAATGATTGAGAGCTTATATGAGCAATTAGTTCAAGAAGATGCGGATGTTTCGAG
CTGTGGTGTGATGAATGTCTATGCTAATGATGAAAGCCACAGTCAGCCAATCAGGATGACTATTTTGT
CTGTGATTCTCAAACATTTCTAAAGGAATACCTCATAGGTGAAAAAATACCTGGGACGATTTGCAATAA
GCTAATCAAGAGACAGATTGCAACTGCCCTATCCTTTCCCTAAGGGGTTGATTTACGAAGATGCCTATTA
CCATTTTGATTTAATCAAGTTGGCCAAGAAAGTATGTGGTTAATACTAAACCCTATTATTACTATTTCCA
TAGAGGGGATAGTATTACGACCAAACCTATGCAGAGAAGGATTTAGCCTATATTGATATCTACCAAAA
GTTTTATAATGAAGTTGTGAAAAACTATCCTGACTTGAAAGAGGTCGCTTTTTTTCAGATTGGCCTATGC
CCACTTCTTTATTTCTGGATAAGATGTTGCTAGATGATCAGTATAAACAGTTTGAAGCCTATTCTCAGAT
TCATCGTTTTTTAAAGGCCATGCCTTTGCTATTTCTAGGAATCCAATTTTCCGTAAGGGGAGAAAGAT
TAGTGCTTTGGCCCTATTCTATAAATATTTCTTATATCGATTCTTATTACTGAAAAATATTGAAAAATC
TAAAAAATTACAT

SP114 amino acid (SEQ ID NO:202)

IQKQTYQNLEIILVDDGATDESGRLCDISIAEQDDRVSVLHKKNEGLSQARNDGMKQAHGDYLI FIDSDD
YIHPEMIQSLYEQLVQEDADVSSCGVMNVYANDESPQSANQDDYFVCDSTFLKEYLIGEKIPGTICNK
LIKROIATALSFPKGLIYEDAYYHFDLIKLAKEYVNTKPYYYYFHRGDSITTKPYAEKDLAYIDIYQK
FYNEVKNYPDLKEVAFFRLAYAHFFILDKMLLDDQYKQFEAYSQIHRFLKGHAFAISRNPFRKGRRI
SALALFINISLYRFLLLKNIKESKKLH

SP115 nucleotide (SEQ ID NO:203)

TAAGGCTGATAATCGTGTTCAAATGAGAACGACGATTAATAATGAATCGCCATTGTTGCTTTCTCCGTT
GTATGGCAATGATAATGGTAACGGATTATGGTGGGGGAACACATTGAAGGGAGCATGGGAAGCTATTCC
TGAAGATGTAAAGCCATATGCAGCGATTGAACTTCATCTGCAAAAGTCTGTAAACCAACAAGTTGTAT
TCCACGAGATACGAAAGAATTGAGAGAATGGTATGTCAAGATGTTGGAGGAAGCTCAAAGTCTAAACAT
TCCAGTTTTCTTGGTTATTATGTGCGGCTGGAGAGCGTAATACAGTTCTCTCAGAGTGGTTAGATGAACA
ATTCCAAAAGTATAGTGTGTTAAAAGGTGTTTTAAATATTGAGAATTATTGGATTTACAATAACCAGTT
AGCTCCGCATAGTGCTAAATATTTGGAAGTTTGTGCCAAATATGGAGCGCATTTTATCTGGCATGATCA
TGAAAAATGGTTCTGGGAAACTATTATGAATGATCCGACATTCTTTGAAGCGAGTCAAAAATATCATAA
AAATTTGGTGTGGCAACTAAAAATACGCCAATAAGAGATGATGCGGGTACAGATTCTATCGTTAGTGG
ATTTTGGTTGAGTGGCTTATGTGATAACTGGGGCTCATCAACAGATACATGGAATGGTGGGAAAAACA
TTATACAAACACATTTGAAACTGGAAGAGCTAGGGATATGAGATCCTATGCATCGGAACCAGAAATCAAT
GATTGCTATGGAATGATGAATGTATATACTGGGGGAGGCACAGTTTATAATTTTGAATGTGCCGCGTA
TACATTTATGACAAATGATGTACCAACTCCAGCATTTACTAAAGGTATTATTCTTTCTTTAGACATGC
TATACAAAATCCAGCTCCAAGTAAGGAAGAAGTTGTAAATAGAACAAAAGCTGTATTTTGAATGGAGA
AGGTAGGATTAGTTCATTAAACGGATTTTATCAAGGACTTTATTTCGAATGATGAAACAATGCCTTTATA
TAATAATGGGAGATATCATATTCTTCTGTAATACATGAGAAAATTGATAAGGAAAAGATTTTCATCTAT

ATTCCCTAATGCAAAAATTTTGACTAAAAATAGTGAGGAATTGTCTAGTAAAGTCAACTATTTAAACTC
GCTTTATCCAAAACCTTTATGAAGGAGATGGGTATGCTCAGCGTGTAGGTAATTCCTGGTATATTTATAA
TAGTAATGCTAATATCAATAAAAAATCAGCAAGTAATGTTGCCTATGTATACTAATAATACAAAAGTCGTT
ATCGTTAGATTTGACGCCACATACTTACGCTGTTGTTAAAGAAAATCCAAATAATTTACATATTTTATT
GAATAATTACAGGACAGATAAGACAGCTATGTGGGCATTATCAGGAAATTTTGATGCATCAAAAAGTTG
GAAGAAAGAAGAATTAGAGTTAGCGAACTGGATAAGCAAAAATTTATCCATCAATCCTGTAGATAATGA
CTTTAGGACAACAACACTTACATTAAGAGGCATACTGGTCATAAACCTCAGATAAATAAGTGGCGA
TAAAAATCATTATACCTTATACAGAAAATTTGGGATGAGAATACCCATGTTTATACCATTACGGTTAATCA
TAATGGAATGGTAGAGATGTCTATAAATACTGAGGGGACAGGTCCAGTCTCTTTCCCAACACCAGATAA
ATTTAATGATGGTAATTTGAATATAGCATATGCAAAACCAACAACACAAAGTTCTGTAGATTACAATGG
AGACCCTAATAGAGCTGTGGATGGTAACAGAAATGGTAATTTTAACTCTGGTTCGGTAACACACACTAG
GGCAGATAATCCCTCTTGGTGGGAAGTCGATTTGAAAAAATGGATAAAGTTGGGCTTGTAAAAATTTA
TAATCGCACAGATGCTGAGACTCAACGCTCTATCTAATTTT

SP115 amino acid (SEQ ID NO:204)

KADNRVQMRTTINNESPLLLSPLYGNDNGNGLWGWNTLKGAWAIPEDVKPYAAIELHPAKVKCKPTSCI
PRDTKELREWYVKMLEEAQSLNIPVFLVIMSAGERNTVPPEWLDEQFQKYSVLKGVLNIENYWIYNNQL
APHSKYLEVCYKGAHFIWHDHEKWFWEITMNDPTFFEASQKYHKNLVLATKNTPIRDDAGTDSIVSG
FWLSGLCDNWGSSTDTWKWWEKHYTNTFETGRARDMSYASEPESMIAMEMMNVTGGGTVYNFECAAY
TFMTNDVPTPAFTKGIIPFFRHAIQNPAPSKEEVNRTKAVFWNGEGRISLNGFYQGLYSNDETMPLY
NNGRYHILPVIHEKIDKEKISSIFPNAKILTKNSEELSSKVNYLNSLYPKLYEGDGYAQRVGNWSYIYN
SNANINKNQVMLPMYTNNTKSLSLDLTPHTYAVVKENPNNLHILLNNYRTDKTAMWALSGNFDASKSW
KKEELELANWISKYNSINPVDNDFRTTTLTLKGHTGHKQPQINISGDKNHYTYTENWDENTHVYITITVNH
NGMVEMSINTEGTGPVSFPTPKDFNDGNLNIAYAKPTTQSSVDYNGDPNRAVDGNRNGNFNSGSVTHTR
ADNPSWWEVDLKKMDKVLVKIYNRTDAETQRLSNF

SP117 nucleotide (SEQ ID NO:205)

CTGTGGCAATCAGTCAGCTGCTTCCAAAACAGTCAGCTTCAGGAACGATTGAGGTGATTTACAGAGAAAA
TGGCTCTGGGACACGGGTGCCTTTCACAGAAATCAGAGGATTCTCAAAAAAGACGGTGATAAAAAAAT
TGACAACACTGCCAAAACAGCTGTGATTCAAAATAGTACAGAAGGTGTTCTCTCAGCAGTTCAAGGGAA
TGCTAATGCTATCGGCTACATCTCCTTGGGATCTTTAACGAAATCTGTCAAGGCTTTAGAGATTGATGG
TGCTCAAGGCTAGTCGAGACACAGTTTATAGATGGTGAATACCCCTCTTCAACGTCCTTCAACATTGTTT
GTCTTCTAATCTTTCCAAGCTAGGTCAAGATTTTATACGCTTTATCCACTCCAACAAGGTCAACAAGT
GGTCACAGATAAATAATTTATTGAAGCTAAAACCGAAACACGGAATATACAAGCCAACACTTATCAGG
CAAGTTGTCTGTTGTAGGTTCCACTTCAGTATCTTCTTTAATGGAAAAATTAGCAGAAGCTTATAAAAA
AGAAAATCCAGAAGTTACGATTGATATTACCTCTAATGGGTCTTCAGCAGGTATTACCGCTGTTAAGGA
GAAAACCGCTGATATTGGTATGGTTTCTAGGGAATTAACCTCTGAAGAAGGTAAGAGTCTCACCCATGA
TGCTATTGCTTTAGACGGTATTGCTGTTGTGGTCAATAATGACAATAAGGCAAGCCAAGTCAGTATGGC
TGAACCTGCAGACGTTTTTATAGTGGCAAATTAACCACCTGGGACAAGATTAAA

SP117 amino acid (SEQ ID NO:206)

CGNQSAASKQSASGTIEVISRENGSGTRGAFTEITGILKKDGDKKIDNTAKTAVIQNSTEGVLSAVQGN
ANAIGYISLGLTKSVKALEIDGVKASRDTVLDGEYPLQRPFNIVWSSNLSKLGQDFISFIHSKQGGQV
VTDNKFIEAKTETTEYTSQHLSGKLSVVGSTSVSSLMEKLAAYKKENPEVTIDITSNGSSAGITAVKE
KTADIGMVSRELTPPEGKSLTHDAIALDGIADVNNNDNKASQVSMAELADVFSGLKTTWDKIK

SP118 nucleotide (SEQ ID NO:207)

TTGTCAACAACAACATGCTACTTCTGAGGGGACGAATCAAAGGCAAAGCAGTTCAGCGAAAGTTCCATG
GAAAGCTTCATACACCAACCTAAACAACCAGGTAAGTACAGAAAGAGGTCAAATCTCTCTTATCAGCTCA
CTTGGATCCAAATGATGTTGATGCATTTTAAATCTCGTTAATGACTATAATACCAATTCGGCTCAAC
TGGCTTATCAGGAGATTTCACTTCTTACTCACACCGAATACGATGTTGAGAAAAATCAGTCACTCTG
GAATCAAAAAGAAGGGCGATTTTGTGGGACCAACTGCCGTATCAATAGTTATTGTCTTTTGAAAAATTC
AGTCACCATTCCAAAGCTTGAAAAAATGACCAGTTGCTTTTCTAGATAATGATGCGATTGATAAAGG
AAAGGTCTTTGATTACAAAGATAAGGAAGAGTTTGATATTCTATTTTCGAGAGTTCCAACCTGAGTCAAC
TACAGATGTCAAGGTTACGCTGAAAAAGATGGAAGCATCTTCTCACAATTTCAATTCAATGAAAAAGC
TCGAATGCTGTCTGTAGTCTTGACGACAATTTGGATGGCGAGTATCTGTTGTAGGCCACGTGGGGT
CTTAGTACCTGCTGATGACGGTTTCTTATTTGTAGAGAAATTGACTTTTGAAGAGCCCTACCAAGCGAT

TAAATTTGCTAGTAAGGAAGATTGCTACAAGTATTTGGGCACCAAGTATGCGGATTATACAGGCGAGGG
ACTGGCTAAGCCTTTTATCATGGATAATGATAAGTGGGTAAACTT

SP118 amino acid (SEQ ID NO:208)

CQQQHATSEGTNQRQSSSAKVPWKASYTNLNNQVSTEEVKSLLSAHLDPNSVDAFFNLVNDYNTIVGST
GLSGDFTSFTHTEYDVEKISHLWNQKKGDFVGTNCRINSYCLLKNSVTIPKLEKNDQLFLDNDIDAIDKG
KVFDSDQKEEFDILFSRVPTESTTDVKVHAEKMEAFFSQFQFNEKARMLSVVLHDNLDGEYLFVGHVGV
LVPADDGFLFVEKLTFEEPYQAIKFASKEDCYKYLGTKYADYTGEGLAKPFI MDNDKWVKL

SP119 nucleotide (SEQ ID NO:209)

TTGTTTCAGGCAAGTCCGTGACTAGTGAACACCAAACGAAAGATGAAATGAAGACGGAGCAGACAGCTAG
TAAAACAAGCGCAGCTAAAGGGAAAGAGGTGGCTGATTTTGAATTGATGGGAGTAGATGGCAAGACCTA
CCGTTTATCTGATTACAAGGGCAAGAAAGTCTATCTCAAATTCTGGGCTTCTTGGTGTTCATCTGTCT
GGCTAGTCTTCCAGATACGGATGAGATTGCTAAAGAAGCTGGTGATGACTATGTGGTCTTGACAGTAGT
GTCACCAGGACATAAGGGAGAGCAATCTGAAGCGGACTTTAAGAATTGGTATAAGGGATTGGATTATAA
AAATCTCCCAGTCTAGTTGACCCATCAGGCAAACCTTTTGAAACTTATGGTGTCCGTTCTTACCCAAC
CCAAGCCTTTATAGACAAAGAAGGCAAGCTGGTCAAACACATCCAGGATTCATGGAAAAAGATGCAAT
TTTGCAAACCTTTGAAGGAATTAGCC

SP119 amino acid (SEQ ID NO:210)

CSGKSVTSEHQTKDEMKEQTASKTSAAKGKEVADFELMGVDGKTYRLSDYKGGKVYLKFWASWCSICL
ASLPDTDEIAKEAGDDYVVLTVVSPGHKGEQSEADFNWYKGLDYKNLPVLVDP SGKLL ETYGVR SYPT
QAFIDKEGKLVKTHPGFMEKDAILQTLKELA

SP120 nucleotide (SEQ ID NO:211)

CTCGCAAATTGAAAAGGCGGCAGTTAGCCAAGGAGGAAAAGCAGTGAAAAAACAGAAATTAGTAAAGA
CGCAGACTTGCACGAAATTTATCTAGCTGGAGGTTGTTTCTGGGGAGTGAGGAATATTTCTCACGTGT
TCCCGGGGTGACGGATGCCGTTTCAGGCTATGCAAATGGTAGAGGAGAAACAACCAAGTACGAATTGAT
TAACCAAACAGGTCATGCAGAAACCGTCCATGTACCTATGATGCCAAGCAAATTTCTCTCAAGGAAAT
CCTGCTTCACTATTTCCGCATTATCAATCCAACCAGCAAAAAATAACAAGGAAATGATGTGGGGACCCA
GTACCGTACTGGTGTATTATTACACAGATGACAAGGATTTGGAAGTGATTAACCAAGTCTTTGATGAGGT
GGCTAAGAAATACGATCAACCTCTAGCAGTTGAAAAGGAAAACCTTGAAGAATTTTGTGGTGGCTGAGGA
TTACCATCAAGACTATCTCAAGAAAAATCCAAATGGCTACTGCCATATCAATGTAAATCAGGCGGCCTA
TCCTGTCTATTGATGCCAGAAATATCCAAAACCAAGTGATGAGGAATTGAAAAAGACCCCTGTCACCTGA
GGAGTATGCAGTTACCCAGGAAAATCAAACAGAACGAGCTTTCTCAAACCGTTACTGGGATAAATTTGA
ATCCGGTATCTATGTGGATATAGCAACTGGGGAACCTCTCTTTTCATCAAAGACAAATTTGAGTCTGG
TTGTGGCTGGCCTAGTTTTACCCAACCCATCAGTCCAGATGTTGTACCTACAAGGAAGATAAGTCCTA
CAATATGACGCGTATGGAAGTGCGGAGCCGAGTAGGAGATTCTCACCTTGGGCATGTCTTTACGGATGG
TCCACAGGACAAGGGCGGCTTACGTTACTGTATCAATAGCCTCTCTATCCGCTTTATTCCCAAAGACCA
AATGGAAGAAAAAGGTACGCTTATTTAC

SP120 amino acid (SEQ ID NO:212)

SQIEKAAVSQGGKAVKKTEISKDADLHEIYLAGGCFWGVVEEYFSRVPGVTDVAVSGYANGRGETTKYELI
NQTGHAETVHVITYDAKQISLKEILLHYFRIINPTSKNKQGNVDVGTQYRTGVYYTDDKDLEVINQVFDEV
AKKYDQPLAVEKENLNKFNVAEDYHQDYLLKNPNGYCHINVNQAAYPVIDASKYPKPSDEELKKTLSP
EYAVTQENQTERAFSNRYWDFESGIYVDIATGEPLFSSKDKFESGCGWPSFTQPI SPDVV TYKEDKSY
NMTRMEVRSRVGDSHLGHVFTDGPQDKGGLRYCINSLSIRFIPKQMEKGTLIY

SP121 nucleotide (SEQ ID NO:213)

TTGTCAAGTCAGGTTCTAATGGTTCTCAGTCTGCTGTGGATGCTATCAAACAAAAAGGGAAATTAGTTGT
GGCAACCAGTCTGACTATGCACCCCTTTGAATTTCAATCATTGGTTGATGGAAAGAACAGGTAGTCGG
TGCAGACATCGACATGGCTCAGGCTATCGCTGATGAACCTGGGGTTAAGTTGGAAATCTCAAGCATGAG
TTTTGACAATGTTTTGACCAGTCTTCAAACCTGGTAAGGCTGACCTAGCAGTTGCAGGAATTAGTGCTAC
TGACGAGAGAAAAGAAGTCTTTGATTTTTCAATCCCATATGAAAACAAGATTAGTTTCTTGGTTTCG
TAAGGCTGATGTGGAATAACAAGGATTTAACTAGCCTAGAAAGTGCTAATATTGCAGCCCAAAAAGG
GACTGTTCCAGAATCAATGGTCAAGGAACAATTGCCAAAAGTTCAATTAACCTCCCTAACTAATATGGG
TGAAGCAGTCAATGAATTGCAGGCTGGAAAAATAGATGCTGTTTCATATGGATGAGCCTGTTGCACTTAG

TTATGCTGCTAAAAACGCTGGCTTAGCTGTCGCAACTGTCAGCTTGAAGATGAAGGACGGCGACGCCAA
TGCC

SP121 amino acid (SEQ ID NO:214)

CQSGSNGSQSAVDAIKQKGLVVATSPDYAPFEFQSLVDGKNQVVGADIDMAQAIADDELGVKLEISSMS
FDNVLTSLOTGKADLAVAGISATDERKEVFDFSIPIYENKISFLVRKADVEKYKDLTSLESANIAAQKG
TVPESMVKEQLPKVQLTSLTNMGEAVNELQAGKIDAVHMDPEVALSYAAKNAGLAVATVSLKMKDGDAN
A

SP122 nucleotide (SEQ ID NO:215)

GGAAACTTCACAGGATTTTAAAGAGAAGAAAACAGCAGTCATTAAGGAAAAAGAAGTTGTTAGTAAAAA
TCCTGTGATAGACAATAACACTAGCAATGAAGAAGCAAAAATCAAAGAAGAAAATTCCAATAAATCCCA
AGGAGATTATACGGACTCATTTGTGAATAAAAAACACAGAAAATCCCAAAAAAGAAGATAAAGTTGTCTA
TATTGCTGAATTTAAAGATAAAGAATCTGGAGAAAAAGCAATCAAGGAACTATCCAGTCTTAAGAATAC
AAAAGTTTTATATACTTATGATAGAATTTTAAACGGTAGTGCCATAGAAACAACCTCCAGATAACTTGGA
CAAAATTAAACAAATAGAAGGTATTTTCATCGGTTGAAAGGGCACAAAAAGTCCAACCCATGATGAATCA
TGCCAGAAAAGGAAATTGGAGTTGAGGAAGCTATTGATTACCTAAAGTCTATCAATGCTCCGTTTGGGAA
AAATTTTGATGGTAGAGGTATGGTCATTTCAAATATCGATACTGGAACAGATTATAGACATAAGGCTAT
GAGAATCGATGATGATGCCAAAGCCTCAATGAGATTTAAAAAAGAAGACTTAAAGGCATGATAAAAA
TTATTGGTTGAGTGATAAAATCCCTCATGCGTTCAATTATTATAATGGTGGCAAAATCACTGTAGAAAA
ATATGATGATGGAAGGGATTATTTTGACCCACATGGGATGCATATTGCAGGGATTCTTGCTGGAAATGA
TACTGAACAAGACATCAAAAACTTTAACGGCATAGATGGAATTGCACCTAATGCACAAATTTTCTCTTA
CAAAATGTATTCTGACGCAGGATCTGGGTTTGCGGGTGATGAAACAATGTTTCATGCTATTGAAGATTC
TATCAACACAACGTTGATGTTGTTTCGGTATCATCTGGTTTTACAGGAACAGGTCTTGAGGTGAGAA
ATATTGGCAAGCTATTCGGGCATTAAGAAAAGCAGGCATTCCAATGGTTGTCGCTACGGGTAACATATGC
GACTTCTGCTTCAAGTTCTTCATGGGATTTAGTAGCAAATAATCATCTGAAAATGACCGACACTGGAAA
TGTAACACGAAGTGCAGCACATGAAGATGCGATAGCGGTGCTTCTGCTAAAAATCAAACAGTTGAGTT
TGATAAAGTTAACATAGGTGGAGAAAAGTTTAAATACAGAAATATAGGGGCCTTTTTCGATAAGAGTAA
AATCACAACAAATGAAGATGGAACAAAAGCTCCTAGTAAATTTAAATTTGTATATATAGGCAAGGGGCA
AGACCAAGATTTGATAGGTTTGGATCTTAGGGGCAAAATTCAGTAATGGATAGAATTTATACAAAGGA
TTTAAAAAATGCTTTTAAAAAAGCTATGGATAAGGGTGCACGCGCCATTATGGTTGTAAATACTGTAA
TTACTACAATAGAGATAATTGGACAGAGCTTCCAGCTATGGGATATGAAGCGGATGAAGGTACTAAAAG
TCAAGTGTTTTCAATTTTCAAGAGATGATGGTGTAAGCTATGGAACATGATTAATCCTGATAAAAAAC
TGAAGTCAAAAGAAATAAAGAAGATTTTAAAGATAAAATTGGAGCAATACTATCCAATGTATATGGA
AAGTTTTTAATTCCAACAAACCGAATGTAGGTGACGAAAAAGAGATTGACTTTAAGTTTGACCTGACAC
AGACAAAGAAGCTCTATAAAGAAGATATCATCGTTTCCAGCAGGATCTACATCTTGGGGGCCAAGAAATAGA
TTTACTTTTAAACCCGATGTTTTCAGCACCTGGTAAAAATATTAAATCCACGCTTAATGTTATTAAATGG
CAAATCAACTTATGGCTATATGTCAGGAAGTAGTATGGCGACTCCAATCGTGGCAGCTTCTACTGTTTTT
GATTAGACCGAAATTAAGGAAATGCTTGAAAGACCTGTATTGAAAAATCTTAAGGGAGATGACAAAAT
AGATCTTACAAGTCTTACAAAAATTGCCCTACAAAATACTGCGCGACCTATGATGGATGCAACTTCTTG
GAAAGAAAAAAGTCAATACTTTGCATCACCTAGACAACAGGGAGCAGGCCTAATTAATGTGGCCAATGC
TTTGAGAAATGAAGTTGTAGCAACTTTCAAAAACACTGATTCTAAAGGTTTGGTAAACTCATATGGTTC
CATTTCTCTTAAAGAAATAAAAGGTGATAAAAAATACTTTACAATCAAGCTTCACAATACATCAAACAG
ACTTTGTACTTTTAAAGTTTTCAGCATCAGCGATAACTACAGATTCTCTAACTGACAGATTAAAACTTGA
TGAAACATATAAAGATGAAAAATCTCCAGATGGTAAGCAAATTTGTTCCAGAAATTCACCCAGAAAAAGT
CAAAGGAGCAAATATCACATTTGAGCATGATACTTTCACTATAGGCGCAAATTTCTAGCTTTGATTTGAA
TGCGGTTATAAATGTTGGAGAGGCCAAAAACAAAAATAAATTTGTAGAATCATTATTTCATTTTGAGTC
AGTGGAAGCGATGGAAGCTCTAAACTCCAGCGGGAAGAAAATAAATTTCCAACCTTCTTTGTCGATGCC
TCTAATGGGATTTGCTGGGAATTGGAACCAACCAATCCTTGATAAATGGGCTTGGGAAGAAGGGTC
AAGATCAAAAACACTGGGAGGTTATGATGATGATGGTAAACCGAAAATTCAGGAACCTTAAATAAGGG
AATTGGTGGAGAACATGGTATAGATAAAATTTAATCCAGCAGGAGTTATACAAAATAGAAAAGATAAAAA
TACAACATCCCTGGATCAAATCCAGAATTTATGCTTTCAATAACGAAGGATCAACGCTCCATCATC
AAGTGGTTCTAAGATTGCTTAACATTTATCCTTTAGATTCAAAATGGAAATCCTCAAGATGCTCAACTTGA
AAGAGGATTAACACCTTCTCCACTTGTATTAAGAAGTGCAGAAGAAGGATTGATT

SP122 amino acid (SEQ ID NO:216)

ETSQDFKEKKTAVIKEKEVSKNPVIDNNTSNEEAKIKEENSNSKSQGDYTDTSFVNKNTENPKKEDKV
VYIAEFKDKESGEKAIKELSSLKNTKVLYTYDRIFNGSAIETTPDNLDKIKQIEGISSVERAQKVQPMNH

ARKEIGVEEAIDYLSINAPFGKNFDGRGMVISNIDTGTDRHKAMRIDDDAKASMRFKKEDLKGTDKN
YWLSDKIPHAFNYNNGGKITVEKYDDGRDYFDPHGMHIAGILAGNDTEQDIKNFNGIDGIAPNAQIFSY
KMYSDAGSGFAGDETMFHAIEDSIKHNVVVSVSSGFTGTGLVGEKYWQAIRALRKAGIPMVVATGNYA
TSASSSSWDLVANNHLKMTDTGNVTRTAAHEDAIIVASAKNQTFEFDKVNIGGESFKYRNIGAFFDKSK
ITTNEGDGTPSKLKFVYIGKGQDQDLIGLDRGKIAVMDRIYTKDLKNAFKKAMDKGARAIMVVNTVN
YYNRDNWTELPMGYEADGTSQVFSISGDDGVKLWNMINPDKKTEVKRNKEDFKDKLEQYYPIDME
SFNSNKNPVGDEKEIDFKFAPDTEKELYKEDIIVPAGSTSWGPRIDLKLPDVSA PGKNIKSTLNVING
KSTYGYMSGTSMATPIVAASTVLIRPKLEMLERPVLKNLKGDDKIDLTSLTKIALQNTARPMMDATSW
KEKSQYFASPRQOGAGLINVANALRNEVVATFKNTDSKGLVNSYGSISLKEIKGDKKYFTIKLHNTSNR
PLTFKVSASAITTDSLTDRLKLDETYKDEKSPDGKQIVPEIHPEKVKGANITFEHDTFTIGANS SFDLN
AVINVGEAKNKNKFVESFIHFESVEAMEALNSSGKKINFQPSLSMPLMGFAGNWNHEPILDKWAWEEGS
RSKTLGGYDDDGKPKIPGTLNKGIGGEHGIDKFNPAQVIQNRKDKNTTSLDQNP ELF AFNNEGINAPSS
SGSKIANIYPLDSNGNPQDAQLERGLTPSPLVLSRAEEGLI

SP123 nucleotide (SEQ ID NO:217)

TGTGGTCGAAGTTGAGACTCCTCAATCAATAACAAATCAGGAGCAAGCTAGGACAGAAAACCAAGTAGT
AGAGACAGAGGAAGCTCCAAAAGAAGACACCTAAAACAGAAAGTCCAAAGGAAGAACCAAAATC
GGAGGTAAAACCTACTGACGACACCTTCCATAAGTAGAAGAGGGGAAAGAAGATTACAGCAGACACAGC
TCCAGTTGAAGAAGTAGGTGAGAGATTGAGTCAAAACAGAGGAAAGTAGCAGTTAAGCCAGAAAG
TCAACCATCAGACAAACCAGCTGAGGAATCAAAAGTTGAACAAGCAGGTGAACCAAGCTCGCGCCAAGAGA
AGACGAAAAGGCACCAGTCGAGCCAGAAAAGCAACCAGAAGCTCCTGAAGAAGAGAAGGCTGTAGAGGA
AACACCGAAAACAAGAAGAGTCAACTCCAGATACCAAGGCTGAAGAACTGTAGAACCAAAAAGAGGAGAC
TGTTAATCAATCTATTGAACAACCAAAAGTTGAAACGCCTGCTGTAGAAAAACAAACAGAACCAACAGA
GGAACCAAAAGTTGAACAAGCAGGTGAACCAAGTCGCGCCAAGAGAAGACGAACAGGCACCAACGGCACC
AGTTGAGCCAGAAAAGCAACCAGAAGTTCCTGAAGAAGAGAAGGCTGTAGAGGAAACACCGAAACCAGA
AGATAAAATAAAGGGTATTGGTACTAAAGAACCAGTTGATAAAAGTGAGTTAAATAATCAAATTGATAA
AGCTAGTTTCAGTTTCTCCTACTGATTATCTACAGCAAGTTACAATGCTCTTGGACCTGTTTTAGAAAC
TGCAAAAGGTGTCTATGCTTCAGAGCCTGTAAAACAGCCTGAGGTAAATAGCGAGACAAATAAACTTAA
AACGGCTATTGACGCTCTAAACGTTGATAAACTGAATTAACAATACGATTGCAGATGCAAAAACAAA
GGTAAAAGAACATTACAGTGATAGAAGTTGGCAAAACCTCCAACTGAAGTTACAAAAGGCTGAAAAGT
TGCAGCTAATACAGATGCTAAACAAAGTGAAGTTAACGAAGCTGTTGAAAAATTAAGTCAACTATTGA
AAAATTGGTTGAATTATCTGAAAAGCCAATATTAACATTGACTAGTACCGATAAGAAAAATATTGGAACG
TGAAGCTGTTGCTAAGTATACTCTAGAAAATCAAAACAAAACAAAATCAAATCAATCACAGCTGAATT
GAAAAAGGAGAAGATTATTAATACTGTAGTCCTTACAGATGACAAGGTAACAACAGAACTATAAG
CGCTGCATTTAAGAAGCTAGAGTACTACAAAGAATACACCCTATCTACAACATGATTTACGACAGAGG
TAACGGTGAAGAACTGAAACTCTAGAAAATCAAAATATTCAATTAGATCTTAAAAAAGTTGAGCTTAA
AAATATTAACGTACAGATTTAATCAAAATACGAAAATGGAAGAACTAATGAATCACTGATAACAAC
TATTCCTGATGATAAGAGCAATTATTATTTAAAAATAACTTCAAATAATCAGAAAACTACATTACTAGC
TGTTAAAAATATAGAAGAACTACGGTTAACGGAACACCTGTATATAAAGTTACAGCAATCGCAGACAA
TTTAGTCTCTAGAACTGCTGATAATAAATTTGAAGAAGAA

SP123 amino acid (SEQ ID NO:218)

VVEVETPQSITNQEQARTENQVVETEAPKEEAPKTEESPKEPKSEVKPTDDTLPKVEEGKEDSAEPA
PVEEVGGEVESKPEEKVAVKPESQPSDKPAEESKVEQAGEPVAPREDEKAPVEPEKQPEAPEEEKAVEE
TPKQEESTPDTKAEETVEPEKEETVNQSI EQPKVETPAVEKQTEPTEEPKVEQAGEPVAPREDEQAPTAP
VEPEKQPEVPEEEKAVEETPKPEDKIKGIGTKPEVDKSELNNQIDKASSVSPTDYSTASYNALGPVLET
AKGVYASEPVKQPEVNSETNKLKTAIDALNVDKTELNNIADAKTKVKEHYSRDSWQNLQTEVTKAQKV
AANTDAKQSEVN EAVEKL TATIEKLVELSEKPILTLTSTDKKILEREAVAKYTLENQNKTKIKSITAE
KKGEEVINTVVLTDKVTETISAAFKNLEYKEYTLSTMIYDRNGEETETLENQNIQLDLKKVELK
NIKRTDLIKYENGKETNESLITTI PDDKSNYYLKITSNNQKTLLAVKNI EETTVNGTPVYKVTAIADN
LVSRTADNKFEET

SP124 amino acid (SEQ ID NO:219)

AACACCTGTATATAAAGTTACAGCAATCGCAGACAATTTAGTCTCTAGAAGTCTGATAATAAATTTGA
AGAAGAATACGTTCACTATATTGAAAAACCTAAAGTCCACGAAGATAATGTATATTATAATTTCAAAGA
ATTAGTGGAAGCTATTCAAAACGATCCTTCAAAAGAATATCGTCTGGGACAATCAATGAGCGCTAGAAA
TGTTGTTCTTAATGGAAAATCATATATCACTAAAGAATTCACAGGAAAACCTTTAAGTTCTGAAGGAAA
ACAATTTGCTATTACTGAATTGGAACATCCATTATTTAATGTGATAACAAACGCAACGATAAATAATGT

GAATTTTGAAGATGTAGAGATAGAACGTTCTGGTCAAGATAATATTGCATCATTAGCCAATACTATGAA
AGGTTCTTCAGTTATTACAAATGTCAAATTACAGGCACACTTTCAGGTCGTAATAATGTTGCTGGATT
TGTAATAATATGAATGATGGAACCTCGTATTGAAAATGTTGCTTCTTTGGCAAACCTACACTCTACAAG
TGGAATGGCTCTCATACAGGGGGAATTGCAGGTACAACTATAGAGGAATTGTTAGAAAAGCATATGT
TGATGCTACTATTACAGGAAACAAAACACGCGCCAGCTTGTAGTTCCTAAAGTAGATTATGGATTAAC
TCTAGACCATCTTATTGGTACAAAAGCTCTCCTAACTGAGTCGGTTGTAAGGTAAAAATAGATGTTTC
AAATCCAGTAGAAGTTGGAGCAATAGCAAGTAAGACTTGGCCTGTAGGTACGGTAAGTAATTCGTGTCAG
CTATGCTAAGATTATCCGTGGAGAGGAGTTATTCGGCTCTAACGACGTTGATGATTCTGATTATGCTAG
TGCTCATATAAAAGATTTATATGCGGTAGAGGGATATTTCGTACGGTAATAGATCATTAGGAAATCTAA
AACATTTACTAAATTAATAAGAACAAAGCTGATGCTAAAGTTACTACTTTCAATATTACTGCTGATAA
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TACAAAAGATTACTTTAACTTGCTTCTAGCAGTGAAGGTTTAAAGTAATGTAAAAGAATATACTATAAC
TGACTTAGGAATTAAATATACACCTAATATCGTTCAAAAAGATAACACTACTCTTGTTAATGATATAAA
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TAGAGTTAATGCAATCAAAGATTATATTTAGAAGAAGCTTCACAGATGTTAAAGAAAACCTAACAAA
CCTAATCACAAAATTAGTTCAAAACGAAGAATCAACTAAATGATTCTCCAGTCTCGTCAAATGAT
TCGTGATAAAGTCGAGAAAAACAAAGCAGCTTTATTACTAGGTTTAACTTACCTAAATCGTTACTATGG
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CGCATTCGGTCAAGTA

SP124 amino acid (SEQ ID NO:220)

TPVYKVTAIADNLVSRADNKFEEYVHYIEKPKVHEDNVYYNFKELVEAIQNDPSKEYRLGQSMSARN
VVPNGKSYITKEFTGKLLSSEGKQFAITELEHPLFNVITNATINNVNFENVEIERSGQDNIALNTMK
GSSVITNVKITGTLSGRNNVAGFVNNMNDGTRIENVAFFGKLHSTSGNGSHTGGIAGTNYRGIVRKAYV
DATITGNKTRASLLVPKVDYGLTLDHLIGTKALLTESVVKGKIDVSNPVEVGAIASKTWPVGTVSNSVS
YAKIIRGEELFGSNDVDDSDYASAHIKDLAYVEGYSSGNRSFRKSKFTFTKLKEQADAKVTTFNITADK
LESDSLPLAKLNEEKAYSSIQDYNAEYNQAYKNLEKLI PFYNKDYIVYQGNKLNKEHHLNLTKEVLSVTA
MNNNEFITNLDEANKIIVHYADGTDYFNLSSESSEGLSNVKEYTITDLGIKYTPNIVQKDNNTLVNDIK
SILESVELQSQTMYQHLNRLGDYRVNAIKDLYLEESFTDVKENLTNLITKLQNEEHQLNDSPAARQMI
RDKVEKNKAALLLGLTYLNRYYGVKFGDVNIKELMLFKPDFYGEKVSVDRLIEIGSKENNIKGSRTFD
AFGQV

SP125 nucleotide (SEQ ID NO:221)

ATTAGACAGATTAATTGAAATCGGTTCTAAAGAGAACAACATTAAAGGTTACGTTACATTCGACGCATT
CGGTCAAGTATTGGCTAAATATACTAAATCAGGTAATTTAGATGCATTTTAAATTATAATAGACAATT
GTTACAAATATAGACAATATGAACGATTGGTTTATTGATGCTACAGAAGACCATGTCTACATCGCAGA
ACGCGCTTCTGAGGTGCAAGAAATTAATAATCTAAACATCGTGCATTTCGATAATTTAAACGAAGTCA
CCTTAGAAATACTATACTCCCACTACTGAATATTGATAAAGCACATCTTTATTTAATTTCAAATTATAA
TGCAATTGCCTTTGGTAGTGACAGACGATTAGGTAAAAAATCATTAGAAGATATTAAAGATATCGTTAA
CAAAGCTGCAGATGGTTATAGAACTATTATGATTTCTGGTATCGTCTAGCGTCTGATAACGTTAAACA
ACGACTACTAAGAGATGCTGTTATTCTTATTTGGGAAGGTTATAACGCTCCTGGTGGATGGGTTGAAAA
ATATGGCCGCTATAATACCGACAAAGTATATACTCCTCTTAGAGAATTCTTTGGTCTCTATGGATAAGTA
TTATAATTATAATGGAACAGGAGCTTATGCTGCTATATATCCTAACTCTGATGATATTAGAACTGATGT
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CAATAAATTAATCAATGGGATAAAATTCGAAATCTAAGTCAAGAAGAGAAAAATGAATTAATTAATCA
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CGAAGCAATTAGCTATAACGATCAATCACCTTATGTAGGTGTTAGAATGATGACCGGTATCTACGGAGG
TAATACTAGTAAAGGTGCTCCTGGAGCTGTTTCATTCAAACATAATGCTTTTAGATTATGGGGTTACTA
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TTCTTCCGTTTTCATCATACGATGATTTACTGACATTGTTTAAAGAAGCTGTTAAAAAAGATGCCGAAAC
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SP125 amino acid (SEQ ID NO:222)

LDRLIEIGSKENNIKGSRTFDAGQVLAKYTKSGNLDNFLNLRQLFTNIDNMNDWFI DATEDHVYIAE
RASEVEEIKNSKHAFDNLKRSHLRNTILPLLNIDKAHLYLISYNAIAFGSAERLGKKSLEDIKDIVN
KAADGYRNYDFWYRLASDNVQRLRLDAVIPIWEGYNAPGGWVEKYGRYNTDKVYTPLEFFGPMDDKY
YNYNGTGAYAAIYPNSDDIRTDVKYVHLEMVGEYGISVYTHETTHVNDRAIYLGFGHREGTDAEAYAQ
GMLQTPVTGSGFDEFGSLGINMVFKRKNDGNQWYITDPKTLKTREDINRYMKGYNDTLTLLDEIEAESV
ISQQNKDLNSAWFKKIDREYRDNNKLNQWDKIRNLSQEEKNELNIQSVNDLVDQQLMTNRNPNNGIYKP
EAI SYNDQSPYVGVRMMGTGIYGGNTSKGAPGAVSFKHNAFRLWGYGYENGFLGYASNKYKQQSKTDGE
SVLSDEYIIKKISNNTFNTEIEFKKAYFKEVKDKATKGLTTFEVNGSSVSSYDDLTLFKEAVKKDAET
LKQEANGNKTVMNNTVKLKEAVYKKLLQQTNSFKTSIFK

SP126 nucleotide (SEQ ID NO:223)

TAAGACAGATGAACGAGCAAGGTGTTGACTTTTCCATTCCCTACTATACTGCAAAAAATAAATCAT
TGTCAAAAATCTGACTTGACTACTTATCAGTCTGTAAACGACTTGGCGCAGAAAAAGGTTGGAGCGCA
GAAAGGTTTCGATTCAAGAGACGATGGCGAAAGATTTGCTACAAAATTCTTCCCTCGTATCTCTGCCTAA
AAATGGGAATTTAATCACAGATTTAAAAATCAGGACAAGTGGATGCCGTTATCTTTGAAGAACCCTGTTTC
CAAGGGATTTGTGGAATAATCCTGATTTAGCAATCGCAGACCTCAATTTTGAAAAAGAGCAAGATGA
TTCCTACGCGGTAGCCATGAAAAAGATAGCAAGAAATTGAAGAGGCAGTTCGATAAAACCATTCAAAA
GTTGAAGGAGTCTGGGGAATTAGACAAACTCATTGAGGAAGCCTTA

SP126 amino acid (SEQ ID NO:224)

KTDERSKVFDFSIPYYTAKNKLIVKKSDDLTYQSVNDLAQKKVGAQKGSIQETMAKDLLQNSSLVSLPK
NGNLITDLKSGQVDAVIFEEPVSKEGFVENNPDLAIADLNFEKEQDDSYAVAMKKDSKKLKRQFDKTIQK
LKESGELDKLIEEAL

SP127 nucleotide (SEQ ID NO:225)

CTGTGAGAATCAAGCTACACCCAAAGAGACTAGCGCTCAAAGACAATCGTCCTTGCTACAGCTGGCGA
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AGATGAAAAACTCAGCGACTACGAGATTCAATTCCAAAGAACCGCCTGGGAGAGCATCTTCCAGGACT
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GAATCAGAAACACACTGATAATCCCGCTACAATTAATTTTTCTGGTGAGGATATTGGTAAACGAATCCT
AGACCTTGCTAACGGAGAGTTTGATTTCCTAGTTTTTGACAAGGTATCCGTTCAAAGATTATCAAGGA
CCGTGGTTTAGACCTCTCAGTCGTTGATTACCTTCTGCAGATAGCCCCAGCAATTATATCATTTTCTC
AAGCGACCAAAAAGAGTTTAAAGAGCAATTTGATAAAGCGCTCAAAGAACTCTATCAAGACGGAACCTT
TGAAAACTCAGCAATACCTATCTAGGTGGTTCTTACCTCCAGATCAATCTCAGTTACAA

SP127 amino acid (SEQ ID NO:226)

CENQATPKETSAQKTIVLATAGDVPPFDYEDKGNLTGFDIEVLKAVDEKLSDYEIQFQRTAWESIFPGL
DSGHYQAAANNLSYTKERAKEYLYSLPISNNPLVLVSNNKNPLTSLDQIAGKTTQEDTGTSNAQFINNW
NQKHTDNPATINFSGEDIGKRILDLANGEFDLFLVFDKVSQKIIKDRGLDLSVVDLPADSPSNYIIFS
SDQKEFKEQFDKALKELYQDGTLEKLSNTYLGGSYLPDQSQLQ

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Table 2
***S. pneumoniae* Antigenic Epitopes**

SP001

Lys-1 to Ile-10; Leu-13 to Lys-32; Arg-41 to Ile-51; Ser-85 to Glu-97; Ala-159 to His-168; Val-309 to Thr-318; Val-341 to Asn-352; Asn-415 to Met-430; Phe-454 to Asn-464; Ser-573 to Gly-591; Asn-597 to Thr-641; and Asn-644 to Ala-664.

SP004

Thr-9 to Thr-24; Ile-29 to Ala-48; Thr-49 to Val-56; Val-286 to Val-312; Pro-316 to Glu-344; Val-345 to Ile-367; Gln-368 to Val-399; Ser-400 to Glu-431; Asn-436 to Ala-457; Ile-467 to Ala-498; and Thr-499 to Glu-540.

SP006

Glu-1 to Lys-13; Pro-24 to Gly-36; Val-104 to Thr-112; Ala-118 to Asn-130; Trp-137 to Ala-146; Ser-151 to Ile-159; Ile-181 to Leu-188; and Pro-194 to Tyr-202.

SP007

Gly-1 to Asn-7; Tyr-24 to Gln-34; His-47 to Phe-55; Ser-60 to Ala-67; Ala-122 to Leu-129; Leu-221 to Lys-230; Val-236 to Phe-256; and Asp-271 to Gly-283; and Leu-291 to Asp-297.

SP008

Leu-4 to Lys-17; Gln-24 to Leu-32; Asp-60 to Ser-66; Ser-70 to Asp-76; Ala-276 to Lys-283; Asn-304 to Lys-311; and Thr-429 to Pro-437.

SP009

Thr-4 to Glu-11; Leu-50 to Asp-60; Ile-102 to Trp-123; and Ser-138 to Ile-157.

SP010

Phe-34 to Gly-41; Asp-44 to Lys-50; Leu-172 to Val-186; Leu-191 to Val-198; Ser-202 to Ile-209; and Val-213 to Leu-221.

SP011

Asn-2 to Thr-10; Asp-87 to Ala-102; Tyr-125 to Glu-132; Thr-181 to Tyr-189; Arg-217 to Thr-232; Asn-257 to Lys-264; Pro-271 to Ser-278; Tyr-317 to Ala-325; Glu-327 to Pro-337; and Thr-374 to Val-381.

SP012

Gly-1 to Lys-19; Phe-34 to Tyr-41; Leu-109 to Lys-126; and Leu-231 to Glu-247.

SP013

Ala-1 to Lys-12; Ile-42 to Pro-53; Leu-138 to Lys-146; Ile-205 to Lys-217; Ser-235 to Ile-251; and Ser-261 to Tyr-272.

SP014

Gly-1 to Val-16; Leu-35 to Leu-44; Asp-73 to Asp-81; Ile-83 to Asp-92; Glu-145 to Ile-153; Phe-188 to Asn-196; Ser-208 to Phe-215; Ile-224 to Leu-231; and Asn-235 to Ala-243.

SP015

Ser-1 to Pro-16; Asn-78 to Glu-88; Ala-100 to Val-108; Ala-122 to Thr-129; Thr-131 to Ser-137; Leu-201 to Ser-220; and Gly-242 to Val-251.

70970

Table 2
***S. pneumoniae* Antigenic Epitopes**

SP016

Gly-1 to Glu-20; Thr-30 to Val-38; Gln-94 to Asn-105; Lys-173 to Pro-182; Gly-189 to Arg-197; Ser-207 to Val-224; Pro-288 to Leu-298; Ala-327 to Ala-342; and Ser-391 to Ala-402.

SP017

Ser-1 to Thr-12; Ala-36 to Tyr-45; Gln-48 to Ile-54; Lys-59 to Lys-76; Tyr-113 to Leu-138; and Phe-212 to Asp-219.

SP019

Val-97 to Glu-117; Asp-163 to Leu-169; Thr-182 to Thr-191; and Lys-241 to Ser-250.

SP020

Asn-18 to Lys-25; Thr-47 to Glu-60; Trp-75 to Val-84; Gly-102 to Val-110; Pro-122 to Ala-131; and Glu-250 to Pro-258.

SP021

Ser-1 to Asp-8; Val-44 to Asp-54; Ala-117 to Val-125; Thr-165 to Thr-173; and Glu-180 to Pro-189.

SP022

Phe-5 to Lys-13; Thr-20 to Ser-36; Glu-59 to Lys-81; Tyr-85 to Gly-93; Trp-94 to Trp-101; and Thr-195 to Trp-208.

SP023

Gln-45 to Glu-59; Asp-69 to Pro-85; Lys-111 to Asn-121; Pro-218 to Ala-228; and Glu-250 to Asn-281.

SP025

Gln-14 to Thr-20; Gly-27 to Phe-33; Gly-63 to Glu-71; and Ile-93 to Phe-102.

SP028

Asp-171 to Pro-179; Tyr-340 to Glu-350; Pro-455 to Tyr-463; and Asp-474 to Pro-480.

SP030

Leu-22 to Leu-37; Trp-81 to Ala-90; Phe-101 to Ala-106; Thr-124 to Tyr-130; and Asn-138 to Glu-144.

SP031

Asp-8 to Val-16; Gly-27 to Thr-35; Gly-178 to Asp-195; Thr-200 to Asp-209; Trp-218 to Leu-224; and Lys-226 to Asp-241.

SP032

Ser-9 to Asp-28; Phe-31 to Val-40; Gly-42 to Arg-50; Ile-52 to Leu-60; Asp-174 to Phe-186; Leu-324 to Met-333; and Thr-340 to Asn-347.

SP033

Gln-2 to Ile-13; Phe-46 to Ile-53; and Asp-104 to Thr-121.

SP034

Glu-36 to Gly-43; Ala-188 to Asp-196; Trp-313 to Gly-320; and Leu-323 to Leu-329.

Table 2
***S. pneumoniae* Antigenic Epitopes**

SP035

Arg-19 to Asp-36; Asp-47 to Val-57; Asn-134 to Thr-143; Asp-187 to Arg-196; and Glu-222 to Ser-230.

SP036

Arg-10 to Arg-17; Lys-29 to Ser-39; Ser-140 to Ala-153; Arg-158 to Tyr-169; Asp-175 to Ala-183; Gly-216 to Asn-236; Ala-261 to Leu-270; Arg-282 to Phe-291; and Thr-297 to Ala-305; Pro-342 to Gln-362; Phe-455 to Asp-463; His-497 to Thr-511; Ala-521 to Gly-529; Ile-537 to Val-546; Ile-556 to Ala-568; Pro-581 to Ser-595; Glu-670 to Ala-685; Ser-696 to Ala-705 and Leu-782 to Ser-791.

SP038

Glu-61 to Pro-69; Phe-107 to Ala-115; Leu-130 to Tyr-141; Ala-229 to Glu-237; Ser-282 to Asn-287; Ala-330 to Glu-338; and Tyr-387 to Glu-393.

SP039

Ser-28 to Asp-35; Pro-88 to Pro-96; Leu-125 to Arg-135; Phe-149 to Leu-157; Gln-246 to Val-254; Ala-357 to Thr-362; Gly-402 to Lys-411; and Leu-440 to Pro-448.

SP040

Thr-21 to Ile-30; His-54 to Gln-68; Arg-103 to Leu-117; and Thr-127 to Leu-136.

SP041

Gly-36 to Asp-49; Leu-121 to Val-128; and Ala-186 to Ile-196.

SP042

Gly-11 to Arg-19; Ile-23 to Lys-31; His-145 to Asn-151; Gln-159 to Asp-166; Ile-175 to Asp-181; Gly-213 to Tyr-225; Ile-283 to Val-291; Pro-329 to Glu-364; Arg-372 to Ser-386; Thr-421 to Phe-430; Leu-445 to Val-453; Ile-486 to Ala-497; Asp-524 to Ala-535; His-662 to Gly-674; and His-679 to Gln-702.

SP043

Lys-2 to Asp-12; Val-58 to Asn-68; Ser-87 to Asp-95; and Asp-102 to Lys-117.

SP044

Gln-3 to Lys-11; Asp-37 to Tyr-52; Glu-171 to Leu-191; His-234 to Asn-247; and Asn-283 to Ala-291.

SP045

Tyr-52 to Ile-63; Asp-212 to Gln-227; Ser-315 to Thr-332; Leu-345 to Phe-354; Asp-362 to Val-370; Thr-518 to Asn-539; Ala-545 to Lys-559; and Val-601 to Pro-610.

SP046

Gln-9 to Ala-18; Glu-179 to Lys-186; Lys-264 to Glu-271; Gly-304 to Glu-17; Ser-503 to Asn-511; Asn-546 to Thr-553; and Asn-584 to Asp-591.

SP048

Table 2
***S. pneumoniae* Antigenic Epitopes**

Tyr-4 to Asp-25; Lys-33 to Val-70; Asp-151 to Thr-170; Asp-222 to Val-257; Thr-290 to Phe-301; and Gly-357 to Val-367.

SP049

Ala-23 to Arg-37; Tyr-85 to Gln-95; Glu-106 to Ile-118; Arg-131 to Ile-144; Gly-150 to Ser-162; and Ala-209 to Asp-218.

SP050

Asp-95 to Glu-113; Gly-220 to Gly-228; Asn-284 to Glu-295; Thr-298 to Val-315.

SP051

Lys-16 to Glu-50; Lys-57 to Asn-104; Ser-158 to Trp-173; Asp-265 to Pro-279; Val-368 to Tyr-386; Glu-420 to Ile-454; Pro-476 to Ile-516; Phe-561 to Gly-581; Thr-606 to Gly-664; and Glu-676 to Val-696.

SP052

Asn-41 to Tyr-60; Phe-80 to Glu-103; Ala-117 to Val-139; Ile-142 to Leu-155; Val-190 to Lys-212; Glu-276 to Phe-283; Arg-290 to Ser-299; Leu-328 to Val-351; Gly-358 to Thr-388; Glu-472 to Ala-483; Val-533 to Asn-561; Asp-595 to Val-606; Glu-609 to Val-620; Glu-672 to Ser-691.

SP053

Ala-62 to Val-101; Thr-147 to Leu-174; Lys-204 to Val-216; Gln-228 to Val-262; Ser-277 to Gly-297; Thr-341 to Gly-368; Thr-385 to Ala-409; Thr-414 to Ser-453; Asn-461 to Leu-490; Glu-576 to Thr-625; Gly-630 to Arg-639; and Asp-720 to Leu-740.

SP054

Glu-7 to Val-28; and Tyr-33 to Glu-44.

SP055

Pro-3 to Val-18; Thr-21 to Lys-53; Val-84 to Lys-99; Ile-162 to Val-172; and Val-204 to Ser-241.

SP056

Val-34 to Tyr-41; Leu-47 to Glu-55; and Pro-57 to Gln-66.

SP057

Asp-1 to Val-25; Pro-29 to Ile-80; Asn-96 to Val-145; and Pro-150 to Glu-172.

SP058

Ala-64 to Thr-70; Leu-82 to His-138; and Val-228 to Asn-236.

SP059

Val-10 to Thr-24; Ser-76 to Pro-102; Ser-109 to Ile-119; Ser-124 to Val-130; Thr-186 to Ile-194; and Asn-234 to Ser-243.

SP060

Leu-70 to Arg-76; and Val-79 to Ile-88.

SP062

Glu-14 to Lys-28; Ser-32 to Lys-46; and Glu-66 to Thr-74.

100

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Table 2

Ile-10 to Val-25; Val-30 to Thr-40; Asp-44 to Pro-54; Asn-57 to Val-63; Pro-71 to Val-100; and Thr-105 to Thr-116.

Pro-12 to Leu-32; Val-40 to Leu-68; Asp-95 to Ala-125; Ser-164 to Glu-184; Ser-314 to Glu-346; Asn-382 to Val-393; Leu-463 to Gln-498; Asn-534 to Lys-548; and Lys-557 to Gly-605.

Asn-2 to Ile-12; Ala-39 to Thr-61; and His-135 to Ala-155.

Gly-1 to Thr-13; Asp-203 to Asn-218; and Gly-240 to Asp-253.

Ser-2 to Ser-12; Val-17 to Gln-26; and Lys-54 to Cys-67.

Ser-32 to Thr-41; Pro-66 to Glu-80; Thr-110 to Val-122; and Val-147 to Thr-180.

Lys-6 to Tyr-16; Gln-19 to Ile-27; Arg-50 to Ala-58; Leu-112 to Val-128; Ile-151 to Asn-167; Leu-305 to Phe-321.

Gln-92 to Asn-158; Gln-171 to Gln-188; Val-204 to Val-240; Thr-247 to Ala-273; Glu-279 to Thr-338; Pro-345 to Glu-368; Asn-483 to Lys-539; Val-552 to Ala-568; Glu-575 to Ser-591; Ser-621 to Gly-640; Gln-742 to Gly-758.

Val-68 to Tyr-81; Tyr-86 to Val-121; Leu-127 to Gly-140; Gly-144 to Ala-155; Gln-168 to Val-185; Asp-210 to Try-241; Glu-246 to Thr-269; Lys-275 to Tyr-295; Gly-303 to Pro-320; Arg-327 to Ile-335; Thr-338 to Thr-364; Tyr-478 to Phe-495; and Tyr-499 to Arg-521.

Glu-37 to Val-45; Glu-55 to Val-68; Thr-104 to Thr-119; Ile-127 to Tyr-135; Asn-220 to Ile-232; Thr-237 to Ala-250; Ser-253 to Ala-263; Glu-284 to Ile-297; and Met-438 to Asn-455.

Gly-2 to Ala-12; Gly-96 to Ile-110; and Thr-220 to Phe-239.

Phe-33 to Tyr-42; Gln-93 to Gly-102; and Val-196 to Asp-211.

Ser-64 to Leu-76; and Phe-81 to Ala-101.

Asp-1 to Glu-12; Tyr-26 to Val-36; and Val-51 to Try-62.

Tabl 2
***S. pneumoniae* Antigenic Epitopes**

SP078

Ala-193 to Ile-208; Tyr-266 to Asn-275; Glu-356 to Leu-369; Ala-411 to Gly-422; Ser-437 to Pro-464; Thr-492 to Glu-534; and Glu-571 to Gln-508.

SP079

Gly-11 to Leu-20; Lys-39 to Leu-48; Leu-72 to Val-85; Asn-147 to Ser-158; Ile-178 to Asp-187; Tyr-189 to Gln-201; and Leu-203 to Ala-216

SP080

Ser-2 to Glu-12; Gln-42 to Ala-51; Ala-116 to Ser-127; Phe-131 to Asp-143; and Ile-159 to Ile-171.

SP081

Gln-2 to Leu-9; Gln-49 to Cys-57; Ile-108 to Val-131; Gly-134 to Leu-145; and Trp-154 to Cys-162.

SP082

Ile-101 to Ser-187; Gly-191 to Asn-221; Arg-225 to Arg-236; Tyr-239 to Leu-255; and Gly-259 to Arg-268.

SP083

Ser-28 to Asp-70.

SP084

Leu-42 to Gln-66; Thr-69 to Lys-81; Glu-83 to Arg-92; and Gly-98 to Asn-110.

SP085

Gln-2 to Val-22; and Ser-45 to Glu-51.

SP086

Leu-18 to Gln-65; and Lys-72 to Val-83.

SP087

Ser-45 to Leu-53; and Thr-55 to Gln-63

SP088

Pro-8 to Ile-16; Leu-25 to Trp-33; Tyr-35 to Gln-43; Leu-51 to Val-59; Val-59 to Arg-67; Thr-55 to Tyr-63; Asn-85 to Gly-93; Thr-107 to Leu-115;
 Leu-115 to Trp-123; Ala-121 to Thr-129; Tyr-153 to Ala-161; His-176 to Gly-184; Tyr-194 to Ala-202; Ala-217 to Gly-225; and Asn-85 to Gly-93.

SP089

Trp-43 to Ala-51; Gln-68 to Phe-76; Val-93 to Gln-101; Phe-106 to Phe-114; Lys-117 to Lys-125; Trp-148 to Phe-156; Glu-168 to Gln-176; Ile-193 to Tyr-201; Lys-203 to Lys-211; Glu-212 to Gln-220; Ile-237 to Tyr-245; Lys-247 to Lys-255; Glu-256 to Gln-264; Met-275 to Gly-283; Lys-286 to Gly-294; Trp-292 to Glu-300; Asp-289 to Thr-297; Tyr-315 to Ser-323; Asp-334 to Lys-342; Pro-371 to Arg-379; Arg-485 to Asn-493; Lys-527 to Arg-535; Phe-537 to Met-545; and Tyr-549 to Glu-557.

SP090

Table 2
***S. pneumoniae* Antigenic Epitopes**

Phe-2 to Gln-10; Gln-13 to Lys-21; Tyr-19 to Glu-27; Tyr-39 to Met-47; Pro-65 to Leu-73; Tyr-121 to His-129; Lys-147 to Ile-155; Gly-161 to Lys-169; Gly-218 to Trp-226; Asp-230 to Thr-238; Tyr-249 to Ala-257; and Ala-272 to Gly-280.

SP091

Ser-19 to Ser-27; Asn-25 to Thr-33; Val-51 to Gln-59; Asn-75 to Asn-83; Ile-103 to Trp-111; Tyr-113 to Ala-121; Leu-175 to Asn-183; Glu-185 to Trp-193; Ala-203 to Tyr-211; Val-250 to Phe-258; Asn-260 to Thr-268; Ser-278 to Asp-286; Tyr-305 to Leu-313; Asn-316 to Gly-324; Asn-374 to Asp-382; Asn-441 to Gly-449; and Ser-454 to Gln-462.

SP092

Arg-95 to Glu-103; Ala-216 to Val-224; Leu-338 to Glu-346; Pro-350 to Ala-358; Pro-359 to Ala-367; Pro-368 to Ala-376; Pro-377 to Ala-385; Pro-386 to Ala-394; Pro-395 to Ala-403; Pro-350 to Ala-358; Gln-414 to Lys-422; Pro-421 to Asn-429; Trp-465 to Tyr-473; Phe-487 to Tyr-495; Asn-517 to Gly-525; Trp-586 to Tyr-594; Phe-608 to Tyr-616; and Asp-630 to Gly-638.

SP093

Gln-30 to Ile-38; Gln-52 to Val-60; Ala-108 to His-116; Tyr-133 to Glu-141; Tyr-192 to Ala-200; and Phe-207 to Ser-215.

SP094

Ala-87 to Val-95; Leu-110 to Cys-118; Gln-133 to Leu-141; Ser-185 to Leu-193; Ile-195 to Gly-203; Asp-206 to Gln-214; Ser-211 to Gly-219; Ile-241 to Thr-249.

SP095

Arg-1 to Gln-9; Phe-7 to Asn-15; Thr-21 to Asn-30; Leu-46 to Phe-54; and Ser-72 to Met-80.

SP096

Gly-29 to Ile-37; Glu-52 to Ser-60; and Leu-64 to Gly-72.

SP097

Ala-11 to Thr-19; Glu-53 to Glu-61; Ser-91 to Lys-99; Thr-123 to Gln-131; and Gly-209 to Lys-217.

SP098

Thr-3 to Ser-11; Gly-38 to Phe-46; Tyr-175 to Asn-183; Met-187 to Cys-195; Gln-197 to Leu-205; Tyr-307 to Gln-315; Gly-318 to Tyr-326; Asn-348 to Val-356; Lys-377 to Pro-385; and Leu-415 to Val-423.

SP099

Arg-19 to Gly-27; Asp-76 to Ser-84; Val-90 to Lys-98; Phe-165 to Val-173; Leu-237 to Pro-245.

SP100

His-111 to Gln-119; Ser-141 to His-149; Asp-154 to Ser-162; Gln-158 to Gln-166; Asp-154 to Gln-166; Lys-180 to Gln-188; and Ser-206 to Gln-214.

SP101

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Table 2
***S. pneumoniae* Antigenic Epitopes**

Glu-23 to Glu-31; Glu-40 to Val-48; Gln-50 to Ser-58; Thr-61 to Ile-69; Leu-82 to Ile-90; Ala-108 to Leu-116; Gln-121 to Pro-129; and Leu-130 to Thr-138.

SP102

Asp-32 to His-40; Arg-48 to Lys-56; and Asp-102 to Thr-110.

SP103

Arg-5 to Gln-13; Gln-22 to Leu-30; Arg-151 to Gln-159; Arg-167 to Gln-175; Pro-189 to Glu-197; Gly-207 to Leu-215; Ser-219 to Gln-227; Ser-233 to Ser-241; Pro-255 to Asp-264; Lys-272 to Gly-280; Ser-318 to Val-326; Thr-341 to Asp-351; Asn-356 to Thr-364; Val-370 to Tyr-378; Ile-379 to Gln-387; and Met-435 to Tyr-443.

SP105

Asn-28 to Pro-36; Thr-77 to Phe-85; Arg-88 to Val-96; Gly-107 to Phe-115; Asp-169 to Asp-177; His-248 to Ser-256; and Ser-274 to Ala-282.

SP106

Val-10 to Thr-18; Ile-62 to Tyr-70; Ile-71 to Pro-79; Lys-86 to Gln-94; Lys-100 to Thr-108; Phe-132 to Leu-140; and Asp-145 to Arg-153.

SP107

Asp-33 to Val-41; and Arg-63 to Gln-71.

SP108

Lys-9 to Gln-17; Leu-44 to Ser-52; Ser-63 to Phe-71; Tyr-109 to Ser-117; Ile-183 to Ile-191; Pro-194 to Leu-202; Gly-257 to Gln-265; Ala-323 to Thr-331; and Leu-381 to Tyr-389.

SP109

Asn-2 to Gln-10; Ala-65 to Lys-73; Leu-76 to Glu-84; Thr-111 to Asp-119; Gln-116 to Tyr-124; Tyr-130 to Val-138; Asp-173 to Gly-181; Asp-196 to Ser-204; Asn-231 to Ser-239; Phe-252 to Ser-260; Phe-270 to Tyr-278; Val-291 to His-299; Asp-306 to Leu-314; and Pro-327 to Gly-335.

SP110

Ser-8 to Glu-16; Ile-37 to Val-45; Ala-107 to Val-115; and Gly-122 to Thr-130.

SP111

Asp-19 to Glu-28; Leu-43 to Ala-51; Asn-102 to Phe-110; Gln-133 to Ser-141; Phe-162 to Asp-170; Tyr-194 to Met-202; and Asp-273 to Ser-281.

Tabl 2

Asp-3 to Gln-11; Gly-21 to Ile-29; Ala-46 to Arg-54; Arg-98 to Arg-106; Thr-114 to Val-122; Gln-133 to Asn-141; and Leu-223 to Thr-231.

Asn-19 to Gly-27; Arg-54 to Ser-62; Val-69 to Gln-77; Ser-117 to Asn-125; Gly-164 to Leu-172; Tyr-193 to Ser-201; Cys-303 to Phe-311; His-315 to Ile-323; Arg-341 to Cys-349; Ile-347 to Ser-355; Arg-403 to Phe-411; Gln-484 to Pro-492; Ser-499 to Leu-507; Ile-541 to Thr-549
Asn-622 to Ile-630; and Glu-645 to Gly-653.

Gly-17 to Leu-25; His-40 to Gln-48; Arg-49 to Arg-57; Ile-65 to Pro-73;
Asn-101 to Asp-111; Gly-128 to Cys-136; Phe-183 to Thr-191; and
Pro-268 to Ile-276.

Met-8 to Ser-16; Tyr-24 to Leu-32; Cys-68 to Leu-76; Ser-100 to Pro-108; Thr-193 to Thr-201; Gly-238 to Pro-250; Thr-280 to Phe-288; Pro-303 to Asn-312; Trp-319 to Leu-328; Leu-335 to Leu-344; Lys-395 to Ala-403; Asn-416 to Gln-424; Tyr-430 to Ser-438; Val-448 to Leu-456; Leu-460 to Thr-468; Pro-502 to Thr-510; Lys-515 to Ile-524; Gln-523 to His-532; Tyr-535 to Thr-543; Ser-559 to Pro-567; Thr-572 to Asn-580; Val-594 to Arg-602; Arg-603 to Asn-611; Thr-620 to Trp-628; and Tyr-644 to Arg-653.

Ala-6 to Gly-14; Ile-19 to Thr-27; Thr-99 to Leu-107; Ser-117 to Asp-125; His-131 to Val-139; Ile-193 to Gly-201; and Val-241 to Gln-249.

Ser-8 to Trp-23; His-46 to Ala-54; Asn-93 to Gly-101; Val-100 to Ser-108; Arg-155 to Asp-163; and His-192 to Leu-200.

Tyr-46 to Lys-54; Ser-93 to Ser-101; Trp-108 to Asn-116; Val-121 to Glu-129; and Tyr-131 to Gln-139.

Ala-57 to Lys-65; Leu-68 to Glu-76; Thr-103 to Tyr-116; Tyr-122 to Val-130; His-163 to Gly-173; Asp-188 to Ser-196; Ser-222 to Ser-231; Phe-244 to Ser-252; Pro-262 to Tyr-270; Val-283 to His-291; and Asp-298 to Leu-306.

Ser-3 to Ala-11; Asp-13 to Leu-21; Ser-36 to Val-44; and Gln-136 to Met-144.

Asn-28 to Lys-36; Glu-39 to Thr-50; Val-54 to Lys-62; Asn-106 to Leu-114; Phe-159 to Gly-167; Asn-172 to Arg-180; Glu-199 to Asn-207;

Table 2
***S. pneumoniae* Antigenic Epitopes**

Lys-230 to His-241; Asn-252 to Gly-263; Met-278 to Ala-287; Thr-346 to Asp-354; Lys-362 to Thr-370; Asp-392 to Asn-405; Asp-411 to Ala-424; Gly-434 to Gly-443; Tyr-484 to Glu-492; Ile-511 to Leu-519; Asn-524 to Asp-538; Glu-552 to Ile-567; Val-605 to Lys-613; Phe-697 to Ala-705; Phe-722 to Leu-730; Leu-753 to Leu-761; Asp-787 to Gln-795; Leu-858 to Asn-866; Ala-892 to Thr-901; Gly-903 to Ile-913; Ile-921 to Asn-931; Asn-938 to Pro-951; Gly-960 to Lys-970; Leu-977 to Asp-985; and Leu-988 to Pro-996.

SP123

Val-4 to Asn-12; Glu-47 to Leu-55; Lys-89 to Glu-100; Ser-165 to Thr-173; Lys-234 to Val-242; Ser-258 to Ser-266; Glu-284 to Asn-292; Tyr-327 to Leu-335; Tyr-457 to Thr-465; Tyr-493 to Glu-501; Thr-506 to Tyr-514; Lys-517 to Thr-525; Asn-532 to Gly-540; and Arg-556 to Glu-564.

SP124

Arg-16 to Glu-24; Gln-52 to Arg-60; Asn-69 to Tyr-77; Glu-121 to Asn-129; Ala-134 to Val-142; Thr-151 to Ala-159; Asn-164 to Glu-172; His-181 to His-189; Thr-210 to Ala-218; Ser-244 to Val-252; Phe-287 to Tyr-297; Ser-312 to Thr-323; His-433 to Tyr-441; Ser-445 to Asn-453; Asn-469 to Thr-477; Asn-501 to Asn-509; Gln-536 to Ala-547; and Gln-608 to Asp-621.

SP125

Ser-9 to Asp-21; Ala-28 to Leu-36; Asn-49 to Phe-57; Val-137 to Arg-145; Asn-155 to Leu-163; Glu-183 to Asp-191; Gly-202 to Tyr-210; Pro-221 to Asp-229; Phe-263 to Ala-271; Phe-300 to Gln-308; Asp-313 to Glu-321; Asn-324 to Asp-332; Ile-346 to Asn-354; Asp-362 to Lys-370; Met-402 to Gly-410; Gly-437 to Gly-445; Ser-471 to Glu-483; Gly-529 to Asp-537; Gln-555 to Val-563; and Leu-579 to Lys-587.

SP126

Leu-22 to Thr-30; Val-65 to Leu-73; and Thr-75 to Asp-83.

SP127

Glu-2 to Ala-12; Asp-28 to Thr-36; Val-105 to Thr-113; Lys-121 to Thr-129; Trp-138 to Pro-146; Ser-152 to Ile-160; Lys-180 to Asp-188; Leu-194 to Asn-202; and Gly-228 to Thr-236.

Table 3
***S. pneumoniae* ORF Cloning Primers**

Primer			
Name	SEQ ID	Sequence	RE
SP001A	NO: 227	GACTGGATCCTAAAATCTACGACAATAAAAAATC	Bam HI
SP001B	NO: 228	CTGAGTCGACTGGTTGTGCTGGTTGAG	Sal I
SP004A	NO: 229	GTCAGGATCCAAATTACAATACGGACTATG	Bam HI
SP004B	NO: 230	CAGTGTGCGACTAAGCTCTAGGTCGGAAAC	Sal I
SP006A	NO: 231	GACTGGATCCTGAGAATCAAGCTACACCCAAAGAG	Bam HI
SP006B	NO: 232	AGTCAAGCTTTTGTAACTGAGATTGATCTGG	Hind III
SP007A	NO: 233	GACTGGATCCTGGTAACCGCTCTTCTCGTAACGCAGC	Bam HI
SP007B	NO: 234	AGTCAAGCTTTTTCAGGAACTTTTACGCTTCC	Hind III
SP008A	NO: 235	AGTCAGATCTTGTGGAAATTTGACAGGTAACAGCAAAAAAGCTGC	Bgl II
SP008B	NO: 236	ACTGAAGCTTTTTTGTGTTTTTCAAGAATTCATCG	Hind III
SP009A	NO: 237	GACTGGATCCTGGTCAAGGAACTGCTTCTAAAGAC	Bam HI
SP009B	NO: 238	AGTCAAGCTTTTCACAAATTCGTTGGTGAAGCC	Hind III
SP010A	NO: 239	GACTGGATCCTAGCTCAGGTGGAAACGCTGGTTCATCC	Bam HI
SP010B	NO: 240	AGTCAAGCTTATCAACTTTTCCACCTTCAACAACC	Hind III
SP011A	NO: 241	GTCAAGATCTCTCCAATATGGTAAATCTGCGGATGG	Bgl II
SP011B	NO: 242	AGTCCTGCAGATCCACATCCGCTTTCATCGGGTTAAAGAAGG	Pst I
SP012A	NO: 243	GACTGGATCCTGGGAAAAATTTCTAGCGAAACTAGTGG	Bam HI
SP012B	NO: 244	GTCACTGCAGCTGTCTCTTCTTACTTCTTTGGTTGC	Pst I
SP013A	NO: 245	GACTGGATCCTGCTAGCGGAAAAAAGATACAACCTTCTGG	Bam HI
SP013B	NO: 246	CTGAAAGCTTTTTTGCCAATCCTTCAGCAATCTTGTC	Hind III
SP014A	NO: 247	GACTAGATCTTGGCTCAAAAAATACAGCTTCAAGTCC	Bgl II
SP014B	NO: 248	AGTCCTGCAGGTTTTTGTGTTGCTTGGTATTGGTCG	Pst I
SP015A	NO: 249	GACTGGATCCTAGTACAACTCAAGCACTAGTCAGACAGAG	Bam HI
SP015B	NO: 250	CAGTCTGCAGTTTCAAAGCTTTTTGTATGTCTTC	Pst I
SP016A	NO: 251	GACTGGATCCTGGCAATTCTGGCGGAAGTAAAGATGC	Bam HI
SP016B	NO: 252	AGTCAAGCTTGTTCATAGCTTTTTTGATTGTTTCG	Hind III
SP017A	NO: 253	GACTGGATCCTTCACAAGAAAAAACAAAAATGAAGATGG	Bam HI
SP017B	NO: 254	AGTCAAGCTTATCGACGTAGTCTCCGCCTTC	Hind III
SP019A	NO: 255	GACTGGATCCGAAAGGTCTGTGGTCAAATAATCTTACC	Bam HI
SP019B	NO: 256	AGTCAAGCTTAGAGTTAACATGGTGCTTGCCAATAGG	Hind III
SP020A	NO: 257	GACTGGATCCAAACTCAGAAAAGAAAGCAGACAATGC	Bam HI
SP020B	NO: 258	AGTCAAGCTTCCAACTGGTTGATCCAAACCATCTG	Hind III
SP021A	NO: 259	GACTGGATCCTTCGAAAGGGTCAGAAGGTGCAGACC	Bam HI
SP021B	NO: 260	AGTCAAGCTTCTGTAGGCTTGGTGTGCCCCAGTTGC	Hind III
SP022A	NO: 261	CTGAGGATCCGGGGATGGCAGCTTTTAAAAATC	Bam HI
SP022B	NO: 262	CAGTAAGCTTGTTTACCCATTACCATTTACC	Hind III
SP023A	NO: 263	CAGTGGATCCAGACGAGCAAAAAATTAAG	Bam HI
SP023B	NO: 264	TCAGAAGCTTGTTTACCCATTACCATTT	Hind III
SP025A	NO: 265	GACTGGATCCCTGTGGTGAGGAAGAACTAAAAAG	Bam HI
SP025B	NO: 266	CTGAGTCGACAATATTCTGTAGGAATGCTTCGAATTTG	Sal I
SP028A	NO: 267	CTGAGGATCCGACTTTTAAACAATAAACTATTGAAGAG	Bam HI
SP028B	NO: 268	GTCAGTGCAGGTTGTACCTCCAAAAATCACGG	Pst I
SP030A	NO: 269	GACTGGATCCCTTTACAGGTAACAACATAAGTCGG	Bam HI
SP030B	NO: 270	CAGTAAGCTTTTTCGAAGTTTGGCTCAGAATTG	Hind III
SP031A	NO: 271	GACTGGATCCCCAGGCTGATACAAGTATCGCA	Bam HI
SP031B	NO: 272	CAGTAAGCTTATCTGCAGTATGGCTAGATGG	Hind III
SP032A	NO: 273	GACTGGATCCGTCTGTATCATTTGAAAAACAAAGAAAC	Bam HI
SP032B	NO: 274	CAGTCTGCAGTTTTTACTGTTGCTGTGCTTGTG	Pst I
SP033A	NO: 275	ACTGAGATCTTGGTCAAAAGGAAAGTCAGACAGGAAAGG	Bgl II
SP033B	NO: 276	CAGTAAGCTTATTCCTGAGCTTTTTTGATAAAGGTTGCGCA	Hind III
SP034A	NO: 277	ACTGGGATCCGAAGGATAGATATATTTTAGCATTTGAGAC	Bam HI
SP034B	NO: 278	AGTCAAGCTTCCATGGTATCAAAGGCAAGACTTGG	Hind III
SP035A	NO: 279	GTCAGGATCCGGTAGTTAAAGTTGGTATTAAACGG	Bam HI
SP035B	NO: 280	AGTCAAGCTTGCAATTTTTGCGAAGTATTCCAAGAG	Hind III
SP036A	NO: 281	AGTCGGATCCTTCTTACGAGTTGGGACTGTATCAAGC	Bam HI

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Table 3
***S. pneumoniae* ORF Cloning Primers**

Primer			
Name	SEQ ID	Sequence	RE
SP036B	NO:282	AGTCAAGCTTGTTTATTTTTCCTTACTTACAGATGAAGG	Hind III
SP038A	NO:283	AGTCGGATCCTACTGAGATGCATCATAATCTAGGAGC	Bam HI
SP038B	NO:284	TCAGCTCGAGTTCTTTGACATCTCCATCATAAGTCGC	Xho I
SP039A	NO:285	GACTGGATCCGGTTTTGAGAAAGTATTTGCAGGGG	Bam HI
SP039B	NO:286	CAGTAAGCTTGGATTTTTTCATGGATGCAATTTTTTTGG	Hind III
SP040A	NO:287	GACTGGATCCGACAACATTTACTATCCATACAGTAGAGTCAGC	Bam HI
SP040B	NO:288	GACTAAGCTTGGCATAAGGTTGCAATTCTGGATTAATTGG	Hind III
SP041A	NO:289	GACTGGATCCGGCTAAGGAAAGAGTGGATG	Bam HI
SP041B	NO:290	GACTAAGCTTTTCATTTTAAATTGACTATGCGCCCCG	Hind III
SP042A	NO:291	GACTGGATCCTTGTTCTATGAACCTGGTTCGTCAAC	Bam HI
SP042B	NO:292	CATGAAGCTTATCCTGGATTTTCCAAGTAAATCT	Hind III
SP043A	NO:293	GACTGGATCCTTATAAGGGTGAATTAGAAAAAGG	Bam HI
SP043B	NO:294	GACTAAGCTTCTTATTAGGATTGTTAGTAGTTG	Hind III
SP044A	NO:295	GACTGGATCCGAATGTTTCAGGCTCAAGAAAGTTCAGG	Bam HI
SP044B	NO:296	GACTAAGCTTTTCCCCTGATGGAGCAAAGTAATACC	Hind III
SP045A	NO:297	GACTGGATCCCTTGGGTGTAACCCATATCCAGCTCCTTCC	Bam HI
SP045B	NO:298	GACTGTGCACTTCAGCTTGTTTATCTGGGGTTGC	Sal I
SP046A	NO:299	GACTGGATCCTAGTGATGCTACTTGGCAAGGAAAACAG	Bam HI
SP046B	NO:300	ACTGCTGCAGATCTTTGCCACCTAGCTTCTCATTG	Pst I
SP048A	NO:301	GTCAGGATCCTGGGATTCAATATGTCAGAGATGATACTAG	Bam HI
SP048B	NO:302	CTAGAAGCTTACGCACCCATTACCATTTATCATTG	Hind III
SP049A	NO:303	GTCAGGATCCGATAATAGAGAAGCATTAATAAAC	Bam HI
SP049B	NO:304	AGTCAAGCTTGACAAAATCTTGAACTCCTCTGGTC	Hind III
SP050A	NO:305	GTCAGGATCCAGATTTTGTGCGAGGAGTGCATACC	Bam HI
SP050B	NO:306	AGTCAAGCTTTCCCTTTTACCCTTACGAATCCAGG	Hind III
SP051A	NO:307	GACTGGATCCATCTGTAGTTTATGCGGATGAAACACTTATTAC	Bam HI
SP051B	NO:308	GACTGTGACGCTTTGGTAGAGATAGAAGTCATG	Sal I
SP052A	NO:309	GACTGGATCCTTACTTTGGTATCGTAGATACAGCCGGC	Bam HI
SP052B	NO:310	AGTCAAGCTTTGTTAATTGCGTACCTTCTAAGCGACC	Hind III
SP053A	NO:311	GACTGGATCCAGCTAAGGTTGCATGGGATGCGATTCC	Bam HI
SP053B	NO:312	GACTGTGACCTGGGCTTTATTAGTTTGAAGTAGC	Sal I
SP054A	NO:313	CAGTGGATCCCTATCACTATGTAAATAAAGAGA	Bam HI
SP054B	NO:314	ACTGAAGCTTTTCTGTCCCTGTTTGAGGCA	Hind III
SP055A	NO:315	CAGTGGATCCTGAGACTCCTCAATCAATAACAAA	Bam HI
SP055B	NO:316	ACGTAAGCTTATAATCAGTAGGAGAACTGAAC	Hind III
SP056A	NO:317	CAGTGGATCCGGATGCTCAAGAACTGCGG	Bam HI
SP056B	NO:318	GACTAAGCTTTTGCCTCTCATTCTTGCTTCC	Hind III
SP057A	NO:319	CAGTGGATCCCGACAAAGGTGAGACTGAG	Bam HI
SP057B	NO:320	ACGTAAGCTTATTTCTTAATTCAAGTGTTTTCTCTG	Hind III
SP058A	NO:321	GACTGGATCCAAATCAATTGGTAGCACAAGATCC	Bam HI
SP058B	NO:322	CAGTGTGACATTAGGAGCCACTGGTCTC	Sal I
SP059A	NO:323	CAGTGGATCCCAAACAGTCAGCTTCAGGAAC	Bam HI
SP059B	NO:324	GACTCTGCAGTTTAATCTTGTCCCAGGTGG	Pst I
SP060A	NO:325	GACTGGATCCATTCGATGATGCGGATGAAAAG	Bam HI
SP060B	NO:326	GACTAAGCTTCATTTGTCTTTGGGTATTTTCGCA	Hind III
SP062A	NO:327	CAGTGGATCCGGAGAGTCGATCAAAAAGTAG	Bam HI
SP062B	NO:328	GTCAGTGCAGTTGCTCGTCTCGAGGTTT	Pst I
SP063A	NO:329	CAGTGGATCCATGGACAACAGGAACTGGGAC	Bam HI
SP063B	NO:330	CAGTAAGCTTATTAGCTTCTGTACCTGTGTTTG	Hind III
SP064A	NO:331	GACTGGATCCCGATGGGCTCAATCCAACCCAGGTCAAGTC	Bam HI
SP064B	NO:332	GACTCTGCAGCATAGCTTTATCCTCTGACATCATCGTATC	Pst I
SP065A	NO:333	GACTGGATCCTTCCAATCAAAAACAGGCAGATGG	Bam HI
SP065B	NO:334	GACTAAGCTTGAGTCCCATAGTCCAAGGCA	Hind III
SP067A	NO:335	AGTCGGATCCTATCACAGGATCGAACGGTAAGACAACC	Bam HI
SP067B	NO:336	ACTGGTGCAGTTCTTTTAACTCCGCTACTGTGTC	Sal I

Table 3
***S. pneumoniae* ORF Cloning Primers**

Primer			
Name	SEQ ID	Sequence	RE
SP068A	NO: 337	CAGTGGATCCAAGTTCATCGAAGATGGTTGGGAAGTCC	Bam HI
SP068B	NO: 338	GATCGTCGACCCGCTCCACATGCTCAACCTT	Sal I
SP069A	NO: 339	TGACGGATCCATCGCTAGCTAGTGAATGCAAGAAAG	Bam HI
SP069B	NO: 340	TGACAAGCTTATTCGTTTTTGAAGTAGTTGCTTTCGT	Hind III
SP070A	NO: 341	GACTGGATCCGACCAGATGGGGCACAAGGTTCAAGG	Bam HI
SP070B	NO: 342	TGACAAGCTTAACTTGTAACGAACAGTTCAATCTG	Hind III
SP071A	NO: 343	GACTAGATCTTTTTTAACCCAACTGTTGGTACTTTCC	Bgl II
SP071B	NO: 344	TGACAAGCTTGTTAGGTGTACATTTTGACCGTC	Hind III
SP072A	NO: 345	ACTGAGATCTTTTTTAACCCAACTGTTGGTACTTTC	Bgl II
SP072B	NO: 346	GACTAAGCTTTCTACGATAACGATCATTTTCTTTACC	Hind III
SP073A	NO: 347	GACTGTGCGACTCGTAGATATTTAAGTCTAAGTGAAGCG	Sal I
SP073B	NO: 348	AGTCAAGCTTGTTAGGTGTTACATTTTGCAAGTC	Hind III
SP074A	NO: 349	GACTGGATCCCTTTGGTTTTGAAGGAAGTAAG	Bam HI
SP074B	NO: 350	TGACCTGCAGACGATTTTTGAAAAATGGAGGTGTATC	Pst I
SP075A	NO: 351	CAGTGGATCCCTACTACCTCTCGAGAGAAAG	Bam HI
SP075B	NO: 352	ACTGAAGCTTTTCGCTTTTACTCGTTTGACA	Hind III
SP076A	NO: 353	CAGTGGATCCTAAGGTCAAAAGTCAGACCGCTAAGAAAGTGC	Bam HI
SP076B	NO: 354	CAGTAAGCTTTAGGGTATCCAAACTGTTGTTGATG	Hind III
SP077A	NO: 355	TGACAGATCTTGACGGGTCTCAGGATCAGACTCAGG	Bgl II
SP077B	NO: 356	TGACAAGCTTCAAAGACATCCACCTCTTGACCTTTG	Hind III
SP078A	NO: 357	GACTGGATCCTAGAGGCTTTGCCAAATGGTGGGAAGGG	Bam HI
SP078B	NO: 358	GTCAGTCGACTTGTTGTAACACTTTTCGAGGTTTGGTACC	Sal I
SP079A	NO: 359	CAGTGGATCCTCAAAAAGAGAAGGAAAACCTGG	Bam HI
SP079B	NO: 360	CAGTCTGCAGTTTCTTCAACAAACCTTGTTCCTG	Pst I
SP080A	NO: 361	CAGTGGATCCACGTTCTATTGAGGACCACTT	Bam HI
SP080B	NO: 362	CAGTAAGCTTTTCCTTCTCAGTCAATTCTTTTCC	Hind III
SP081A	NO: 363	GACTGGATCCCGCTCAAAATACCAGAGGTGTTTCAAG	Bam HI
SP081B	NO: 364	GACTAAGCTTAGTACCATGGGTGTGACAGGTTTGAA	Hind III
SP082A	NO: 365	CTGAGGATCCAATTGTACAATTAGAAAAAGATAGC	Bam HI
SP082B	NO: 366	TGACAAGCTTGCGTTGACTAGGTTCTGCAATGCC	Hind III
SP083A	NO: 367	GACTGGATCCTCTGACCAAGCAAAAAGAGCAGTCAATGA	Bam HI
SP083B	NO: 368	TCAGCAGCTGATCATTGACTTTACGATTTGCTCC	Bgl II
SP084A	NO: 369	GACTGGATCCGTCCGGCTCTGTCAGTCCACTTTTTCAGCG	Bam HI
SP084B	NO: 370	TCAGAAAGCTTATTTTCTTTTCCCTTAATGCGTT	Hind III
SP085A	NO: 371	GACTGGATCCGGGACAAATTCAAAAAAATAGGCAAGAGG	Bam HI
SP085B	NO: 372	GTCAAAGCTTTGGCTCTTTGATTGCCAACAACTG	Hind III
SP086A	NO: 373	GACTGGATCCTCGCTACCAGCAACAAAGCGAGCAAAAGG	Bam HI
SP086B	NO: 374	GACTAAGCTTACTTTTTTCTTTTTTCCACACGA	Hind III
SP087A	NO: 375	CAGTGGATCCGAACCGACAAGTCGCCCACATCAAGACT	Bam HI
SP087B	NO: 376	CTGAAAGCTTTGAATTCTCTTTCTTTTCAAGGCT	Hind III
SP088A	NO: 377	TCGAGGATCCGGTTGTGCGCTGGCAATATATCCCGT	Bam HI
SP088B	NO: 378	CAGTAAGCTTCCGAACCCATTGCGCATTTATAGTTGAC	Hind III
SP089A	NO: 379	AGTCGGATCCGGCCAAATCAGAATGGGTAGAAGAC	Bam HI
SP089B	NO: 380	TGACCTGCAGCTTCTCATTTGATTTTCATCATCAC	Pst I
SP090A	NO: 381	GACTGGATCCATTTGCAGATGATTCTGAAGGATGG	Bam HI
SP090B	NO: 382	TCAGCTGCAGCTTAACCCATTACCATTTCTAGTTTAAG	Pst I
SP091A	NO: 383	GACTGGATCCTGTGCTGCTGCAAAATGAACTGAAGTAGC	Bam HI
SP091B	NO: 384	GACTAAGCTTATACCAAACGCTGACATCTACGCG	Hind III
SP092A	NO: 385	AGTCAGATCTTACGTCTCAGCCTACTTTTGTAAGAGC	Bgl II
SP092B	NO: 386	GACTAAGCTTAACCCATTACCATTTGGCATTGAC	Hind III
SP093A	NO: 387	CAGTGGATCCTGGACAGGTGAAAGGTCATGCTACATTTGTG	Bam HI
SP093B	NO: 388	GACTAAGCTTCAACCATTTGAGACCTTGCAACAC	Hind III
SP094A	NO: 389	GTCAGGATCCGATTGCTCCTTTGAAGGATTTGAGAGAAACC	Bam HI
SP094B	NO: 390	GACTAAGCTTCGATCAAAGATAAGATAAATATATATAAAGT	Hind III
SP095A	NO: 391	GACTGGATCCTAGGTCATATGGGACTTTTTTCTACAACAAAATAGG	Bam HI

Table 3
***S. pneumoniae* ORF Cloning Primers**

Primer			
Name	SEQ ID	Sequence	RE
SP095B	NO: 392	TGACAAGCTTATCTATCAGCTCATTTAATCGTTTTTG	Hind III
SP096A	NO: 393	CTGAGGATCCCAACGTTGAGAATTATTTGCGAATG	Bam HI
SP096B	NO: 394	TGACAAGCTTGAGTCTACAAAAGTAATGTAC	Hind III
SP097A	NO: 395	GTCAGGATCCCTACTATCAATCAAGTTCTTCAGCC	Bam HI
SP097B	NO: 396	TGACAAGCTTGACTGAGGCTTGGACCAGATTGAAAAG	Hind III
SP098A	NO: 397	GACTGGATCCGACAAAAACATTAAAAACGTCCTGAGG	Bam HI
SP098B	NO: 398	GACTAAGCTTAGCACGAAGTGTGACGCTGGTTCC	Hind III
SP099A	NO: 399	GACTGGATCCTTCTCAGGAGACCTTTAAAAATATC	Bam HI
SP099B	NO: 400	GACTAAGCTTGTTGGCCATCTTGTACATACC	Hind III
SP100A	NO: 401	GACTGGATCCAGTAAATGCGCAATCAAATTC	Bam HI
SP100B	NO: 402	AGTCCTGCAGGTATTTAGCCCCAATAATCTATAAAGCT	Pst I
SP101A	NO: 403	CAGTGGATCCTTACCGCGTTTCATCAAGATGTC	Bam HI
SP101B	NO: 404	GACTAAGCTTGCCAGATGTTGAAAAGAGAGTG	Hind III
SP102A	NO: 405	GACTGGATCCGTGGATGGGCTTTAACTATCTTCGTATTCG	Bam HI
SP102B	NO: 406	AGTCAAGCTTGCTAGTCTTCACTTTCCCTTTCC	Hind III
SP103A	NO: 407	GACTGTCGACACTAAACCAGCATCGTTCGCAGGA	Sal I
SP103B	NO: 408	CTGACTGCAGCTTCTTGAAGAAATAATGATTGTGG	Pst I
SP105A	NO: 409	CAGTGGATCCTGACTACCTTGAAATCCCACTT	Bam HI
SP105B	NO: 410	CAGTAAGCTTTTTTTTTTAAAGTTGTAGAATGATTTCAATC	Hind III
SP106A	NO: 411	CAGTGTGCGACTCGTATCTTTTTTTGGAGCAATGTT	Sal I
SP106B	NO: 412	GACTAAGCTTAAATGTTCCGATACGGGTGATTG	Hind III
SP107A	NO: 413	CAGTGGATCCGGAATCTCTCAAAGATGTGAAAG	Bam HI
SP107B	NO: 414	GACTAAGCTTCTTGAGTTTGTCAAGGATTGCTTT	Hind III
SP108A	NO: 415	CAGTGGATCCCAAGAAATCCTATCATCTCTTCCAGAAG	Bam HI
SP108B	NO: 416	GACTAAGCTTTTTCAGAACTAAAAGCCGCAGCTT	Hind III
SP109A	NO: 417	GACTGGATCCACGAAATGCAGGGCAGACAG	Bam HI
SP109B	NO: 418	CAGTAAGCTTATCAACATAATCTAGTAAATAAGCGT	Hind III
SP110A	NO: 419	CAGTGGATCCTGTATAGTTTTTTAGCGCTTGTTCTTC	Bam HI
SP110B	NO: 420	GTCAAAGCTTTGATAGAGTGTCAATAATCTTCTTTAG	Hind III
SP111A	NO: 421	GACTGGATCCGTGTGTGCGAGCATATTCTGAAG	Bam HI
SP111B	NO: 422	CAGTAAGCTTACTTTTACCATTCTTTGTCTGCATC	Hind III
SP112A	NO: 423	GACTGTCGACGTGTTTGGATAGCATTCAGAATCAGACG	Sal I
SP112B	NO: 424	CAGTAAGCTTCGGAAGTAAAGACAATTTTTTCC	Hind III
SP113A	NO: 425	CAGTGGATCCGTGCCTAGATAGTATTACTCAAAC	Bam HI
SP113B	NO: 426	GACTAAGCTTTTTTGCTTATTTCTCTCAATTTTTTC	Hind III
SP114A	NO: 427	CAGTGGATCCCATTCAAGAGCAGACCTATCAAAATC	Bam HI
SP114B	NO: 428	ACTGAAGCTTATGTAATTTTTTTAGATTTTTTCAATTTTTTCAG	Hind III
SP115A	NO: 429	AGTCGGATCCTAAGGCTGATAATCGTGTTCAAATG	Bam HI
SP115B	NO: 430	GACTAAGCTTAAAATTAGATAGACGTTGAGT	Hind III
SP117A	NO: 431	AGTCGGATCCCTGTGGCAATCAGTCAGCTGCTTCC	Bam HI
SP117B	NO: 432	GACTGTCGACTTTAATCTTGTCCAGGTGGTTAATTTGCC	Sal I
SP118A	NO: 433	ACTGGTCGACTTGTCAACAACAACATGCTACTTCTGAG	Sal I
SP118B	NO: 434	GACTCTGCAGAAGTTTAACCCACTTATCATTATCC	Pst I
SP119A	NO: 435	ACTGGGATCCTTGTTTCAGGCAAGTCCGTGACTAGTGAAC	Bam HI
SP119B	NO: 436	GACTAAGCTTGGCTAATTCCTTCAAAGTTTGCA	Hind III
SP120A	NO: 437	AGTCGGATCCCTCGCAAATTGAAAAGGCGGCAGTTAGCC	Bam HI
SP120B	NO: 438	GACTAAGCTTGTAATAAGCGTACCTTTTTTCTTCC	Hind III
SP121A	NO: 439	TCAGGGATCCTTGTCAGTCAGGTTCTAATGGTTCTCAG	Bam HI
SP121B	NO: 440	AGTCAAGCTTGGCATTGGCGTCGCCGTCCTTC	Hind III
SP122A	NO: 441	GACTGGATCCGGAACCTTACAGGATTTTAAAGAGAAG	Bam HI
SP122B	NO: 442	GACTGTCGACAATCAACTTCTTCTGCACTTCT	Sal I
SP123A	NO: 443	CAGTGGATCCTGTGGTCAAGTTGTGAGACTCCTCAATC	Bam HI
SP123B	NO: 444	GACTAAGCTTTTTCTTCAAATTTATTATCAGC	Hind III
SP124A	NO: 445	AGTCGGATCCAACACCTGTATATAAAGTTACAGCAATCG	Bam HI
SP124B	NO: 446	GACTGTCGACTACTTGACCGAATGCGTCGAATGTACG	Sal I